

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
15 November 2001 (15.11.2001)

PCT

(10) International Publication Number
WO 01/85772 A2

(51) International Patent Classification⁷: C07K 14/195

(21) International Application Number: PCT/GB01/02003

(22) International Filing Date: 8 May 2001 (08.05.2001)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:
0011108.8 8 May 2000 (08.05.2000) GB

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(81) Designated States (national): AE, AG, AL, AM, AT, AU,
AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU,
CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH,
GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC,
LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW,
MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK,
SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA,
ZW.

(84) Designated States (regional): ARIPO patent (GH, GM,
KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian
patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European
patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE,
IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF,
CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).

Published:

— without international search report and to be republished
upon receipt of that report

For two-letter codes and other abbreviations, refer to the "Guid-
ance Notes on Codes and Abbreviations" appearing at the begin-
ning of each regular issue of the PCT Gazette.



WO 01/85772 A2

(54) Title: VIRULENCE GENES, PROTEINS, AND THEIR USE

(57) Abstract: A series of genes from *Neisseria meningitidis* are shown to encode products which are implicated in virulence. The identification of these genes therefore allows attenuated microorganisms to be produced. Furthermore, the genes or their encoded products can be used in the manufacture of vaccines for therapeutic application.

VIRULENCE GENES, PROTEINS, AND THEIR USE

Field of the Invention

This invention relates to virulence genes and proteins, and their use. More particularly, it relates to genes and proteins/peptides obtained from *Neisseria meningitidis*, and their use in therapy and in screening for drugs.

Background of the Invention

Neisseria meningitidis is a Gram-negative bacterial pathogen that is implicated in septic shock and bacterial meningitis. This bacterium is a leading cause of bacterial meningitis in developed countries, and causes large-scale epidemics in Africa and China. In the UK, *Neisseria meningitidis* is the leading cause of death in childhood apart from road traffic accidents. The bacterium naturally inhabits the human nasopharynx and then gains access to the blood stream from where it causes severe septicaemia or meningitis. Although current anti-microbials are effective in eliminating the bacterium from the body, the mortality from meningococcal septicaemia remains substantial. It would be desirable to provide means for treating or preventing conditions caused by *Neisseria meningitidis*, e.g. by immunisation.

Summary of the Invention

The present invention is based on the discovery of virulence genes in *Neisseria meningitidis*.

According to a first aspect of the invention, a peptide of the invention is encoded by an operon including any of the nucleotide sequences identified in claim 1, or a homologue thereof in a Gram-negative bacterium, or a functional fragment thereof, for therapeutic or diagnostic use.

The peptides may have many therapeutic uses for treating *Neisseria* infections, including use in vaccines for prophylactic application.

According to a second aspect, a polynucleotide encoding a peptide defined above, may also be useful for therapy or diagnosis.

According to a third aspect, the genes that encode the peptides may be utilised to prepare attenuated microorganisms. The attenuated microorganisms will usually have a mutation that disrupts the expression of one or more of the genes identified herein, to provide a strain that lacks virulence. These microorganisms will also have use in therapy and diagnosis.

According to a fourth aspect, the peptides, genes and attenuated microorganisms according to the invention may be used in the treatment or prevention of a condition associated with infection by *Neisseria* or Gram-negative bacteria.

Description of the Invention

The present invention is based on the discovery of genes encoding peptides which are implicated in virulence. The peptides and genes of the invention are therefore useful for the preparation of therapeutic agents to treat infection. It should
5 be understood that references to therapy also include preventative treatments, e.g. vaccination. Furthermore, while the products of the invention are intended primarily for treatment of infections in human patients, veterinary applications are also considered to be within the scope of the invention.

The present invention is described with reference to *Neisseria meningitidis*.
10 However, all the *Neisseria* strains, and many other Gram-negative bacterial strains are likely to include related peptides or proteins having amino acid sequence identity or similarity to those identified herein. Organisms likely to contain the peptides include, but are not limited to the genera *Salmonella*, *Enterobacter*, *Klebsiella*, *Shigella* and *Yersinia*.

15 The experiments carried out to identify the virulence genes of the invention utilised *N. meningitidis* strain B. Homology searches were performed on the strain A database, however the proteins and genes from strain B are preferred.

Preferably, the peptides that may be useful in the various aspects of the invention have greater than a 40% similarity with the peptides identified herein. More
20 preferably, the peptides have greater than 60% sequence similarity. Most preferably, the peptides have greater than 80% sequence similarity, e.g. 95% similarity. With regard to the polynucleotide sequences identified herein, related polynucleotides that may be useful in the various aspects of the invention may have greater than 40% identity with the sequences identified herein. More preferably, the polynucleotide
25 sequences have greater than 60% sequence identity. Most preferably, the polynucleotide sequences have greater than 80% sequence identity, e.g. 95% identity.

The terms "similarity" and "identity" are known in the art. The use of the term "identity" refers to a sequence comparison based on identical matches between correspondingly identical positions in the sequences being compared. The term
30 "similarity" refers to a comparison between amino acid sequences, and takes into account not only identical amino acids in corresponding positions, but also functionally similar amino acids in corresponding positions. Thus similarity between polypeptide sequences indicates functional similarity, in addition to sequence similarity.

Levels of identity between gene sequences and levels of identity or similarity
35 between amino acid sequences can be calculated using known methods. In relation

to the present invention, publicly available computer based methods for determining identity and similarity include the BLASTP, BLASTN and FASTA (Atschul *et al.*, J. Molec. Biol., 1990; 215:403-410), the BLASTX program available from NCBI, and the Gap program from Genetics Computer Group, Madison WI. The levels of similarity and identity provided herein, were obtained using the Gap program, with a Gap penalty of 12 and a Gap length penalty of 4 for determining the amino acid sequence comparisons, and a Gap penalty of 50 and a Gap length penalty of 3 for the polynucleotide sequence comparisons.

Having characterised a gene according to the invention, it is possible to use the gene sequence to search for related genes or peptides in other microorganisms. This may be carried out by searching in existing databases, e.g. EMBL or GenBank.

Peptides or proteins according to the invention may be purified and isolated by methods known in the art. In particular, having identified the gene sequence, it will be possible to use recombinant techniques to express the genes in a suitable host. Active fragments and related molecules can be identified and may be useful in therapy. For example, the peptides or their active fragments may be used as antigenic determinants in a vaccine, to elicit an immune response. They may also be used in the preparation of antibodies, for passive immunisation, or diagnostic applications. Suitable antibodies include monoclonal antibodies, or fragments thereof, including single chain Fv fragments. Methods for the preparation of antibodies will be apparent to those skilled in the art.

Active fragments of the peptides are those that retain the biological function of the peptide. For example, when used to elicit an immune response, the fragment will be of sufficient size, such that antibodies generated from the fragment will discriminate between that peptide and other peptides on the bacterial microorganism. Typically, the fragment will be at least 30 nucleotides (10 amino acids) in size, preferably 60 nucleotides (20 amino acids) and most preferably greater than 90 nucleotides (30 amino acids) in size.

It should also be understood, that in addition to related molecules from other microorganisms, the invention encompasses modifications made to the peptides and polynucleotides identified herein which do not significantly alter the biological function. It will be apparent to the skilled person that the degeneracy of the genetic code can result in polynucleotides with minor base changes from those specified herein, but which nevertheless encode the same peptides. Complementary polynucleotides are also within the invention. Conservative replacements at the amino acid level are also

envisaged, i.e. different acidic or basic amino acids may be substituted without substantial loss of function.

The preparation of vaccines based on attenuated microorganisms is known to those skilled in the art. Vaccine compositions can be formulated with suitable carriers or adjuvants, e.g. alum, as necessary or desired, to provide effective immunisation against infection. The preparation of vaccine formulations will be apparent to the skilled person. The attenuated microorganisms may be prepared with a mutation that disrupts the expression of any of the genes identified herein. The skilled person will be aware of methods for disrupting expression of particular genes. Techniques that may be used include insertional inactivation or gene deletion techniques. Attenuated microorganisms according to the invention may also comprise additional mutations in other genes, for example in a second gene identified herein or in a separate gene required for growth of the microorganism, e.g. an *aro* mutation or, with regard to *Salmonella*, in a gene located in the SPI2 region identified in WO-A-96/17951.

Attenuated microorganisms may also be used as carrier systems for the delivery of heterologous antigens, therapeutic proteins or nucleic acids (DNA or RNA). In this embodiment, the attenuated microorganisms are used to deliver a heterologous antigen, protein or nucleic acid to a particular site *in vivo*. Introduction of a heterologous antigen, peptide or nucleic acid into an attenuated microorganism can be carried out by conventional techniques, including the use of recombinant constructs, e.g. vectors, which comprise polynucleotides that express the heterologous antigen or therapeutic protein, and also include suitable promoter sequences. Alternatively, the gene that encodes the heterologous antigen or protein may be incorporated into the genome of the organism and the endogenous promoters used to control expression.

More generally, and as is well known to those skilled in the art, a suitable amount of an active component of the invention can be selected, for therapeutic use, as can suitable carriers or excipients, and routes of administration. These factors would be chosen or determined according to known criteria such as the nature/severity of the condition to be treated, the type and/or health of the subject etc.

In a separate embodiment, the products of the invention may be used in screening assays for the identification of potential antimicrobial drugs or for the detection for virulence. Routine screening assays are known to those skilled in the art, and can be adapted using the products of the invention in the appropriate way. For example, the products of the invention may be used as the target for a potential drug,

with the ability of the drug to inactivate or bind to the target indicating its potential antimicrobial activity.

The various products of the invention may also be used in veterinary applications.

5 The following is a brief overview of the experimental procedure used to identify the virulence genes.

Signature-tagged mutagenesis (STM) (Hensel *et al.*, Science, 1995; 269: 400-403) was used to identify genes in *N. meningitidis* that are essential for septicemic infection. In STM, individual mutants are tagged with unique sequence identifiers, allowing large numbers of mutants to be analyzed simultaneously. However, it is necessary to construct libraries of insertional mutants, so far a limitation in studying *N. meningitidis*. Mutagenesis was accomplished successfully using a method in which *Neisseria* DNA is modified *in vitro* using purified components of Tn10 transposition. As *N. meningitidis* efficiently takes up exogenous DNA, the modified alleles are then introduced into *N. meningitidis* by transformation. The mutants can then be screened for their ability to cause systemic infection.

The vector pSTM115 (Sun *et al.* Nature Medicine, 2000; 6(11): 1269-1273) was used as the transposon donor for *in vitro* mutagenesis. 96 pSTM115 derivatives, each containing unique signature tags, were included in 96 separate transposition reactions. The modified genomic DNA was repaired, and returned to the host by transformation. To determine whether Tn10 insertion occurs at diverse sites, 40 transformants were assessed from a single transposition reaction by Southern blot analysis. Each had a single, distinct Tn10 insertion. To establish whether Tn10 integration was stable during systemic infection of infant rats, the hybridization patterns of six mutants before and after passage through rats were compared. Identical hybridization patterns before and after infection were obtained.

The experimental conditions used were as follows:

Bacterial strains and growth:

C311+ is an ET-5, serogroup B *N. meningitidis* isolate from a patient with invasive meningococcal infection (Virji *et al.*, Mol. Microbiol., 1991; 5: 1831-1841). *N. meningitidis* was grown on brain-heart infusion medium with 5% Levinthal's supplement. *E. coli* strains were propagated on Luria Bertani media. Kanamycin was added to solid media as required at concentrations of 75 and 50 µg/ml for *N. meningitidis* and *E. coli*, respectively.

35 *In vitro* transposition and insertion site characterization:

The pACYC184 origin of replication in pSTM115 was used to isolate the insertion site by marker rescue. Nucleotide sequencing was carried out using the dye-termination method (Perkin Elmer, Norwalk, Connecticut) with primers NG62 (5'-TTGGTTAATTGGTTGTAACACTGG-3') (SEQ ID NO. 209) or NG99 (5'-ATTCTCATGTTTGACAGCG-3') (SEQ ID NO. 210). Homology searches were performed against protein databases (<http://www.ncbi.nlm.nih.gov/>), including the serogroup A and B *N.meningitidis* and the *N. gonorrhoeae* genome sequences (http://www.sanger.ac.uk/Projects/N_meningitidis and <http://www.tigr.org>, and <http://dna1.chem.ou.edu/gono.html>, respectively).

10 Tag amplification, cloning and Southern blot analyses:

Hybridizations and preparations of dot blots were performed as described in Hensel *et al.*, *supra*, except that tags were amplified with primers NG13 (5'-ATCCTACAACCTCAAGCT-3') (SEQ ID NO. 211) and NG14 (5'-ATCCATTCTAACCAAGC-3') (SEQ ID NO. 212), and PCR products, rather than plasmid DNA, were fixed onto membranes. Oligonucleotides S1 (5'-AAGAGATTACGCGCAGACC-3') (SEQ ID NO. 213) and S2 (5'-AATACGCAACCGCCTCTC-3') (SEQ ID NO. 214) anneal to sequences in pSTM115 flanking the 'signature tags' and were used to amplify a 367-base-pair product from each pSTM115 derivative. For Southern blot analysis, the kanamycin-resistance cassette from pSTM115 was labelled using the random primers method (NEB), and was used as a probe against genomic DNA digested with *Cla*I.

Animal model:

For screening the STM pools, mutants were grown individually for 18 h in microtiter plates. The bacteria were pooled, then re-suspended in PBS. Wistar rats (5 days old) were inoculated intraperitoneally with 100 µl of the suspension, and were monitored for 48 h. To establish the competitive index of a mutant, wild-type and mutant bacteria were grown for 18 h on solid media and collected into PBS, and rats were inoculated with a 1:1 ratio of mutant to wild-type cells in a total inoculum of 5×10^6 CFU. The proportion of mutant (kanamycin-resistant) to wild-type (kanamycin-sensitive) bacteria was determined by plating replicate samples to media with or without added antibiotic.

The results of the homology searches are shown in Table 1.

Table 1

SEQ ID NO.	PROTEIN	ACCESSION NO.	ORGANISM
3 & 4	3'dehydroquinate synthase	NMB0647	N. gonorrhoeae
25 & 26	Glycosyl transferase	LSI2	N. gonorrhoeae
35 & 36	Polyribonucleotide nucleotidyl transferase	NMB0758	N. meningitidis
37 & 38	Phosphoribosyl formyl glycinamide synthase	PURL	E. coli
43 & 44	Shikimate dehydrogenase	AROE	N. meningitidis
47 & 48	Hypothetical 21.7 kD protein	NMB0673	N. meningitidis
53 & 54	Putative cell binding factor	NMB0345	N. meningitidis
57 & 58	Hypothetical protein	HI0633	H. influenzae
61 & 62	Na ⁺ /H ⁺ antiporter	NMBN0536	N. meningitidis
63 & 64	Chorismate synthase	AROC	V. anguillarum
67 & 68	Paraquat-inducible protein B	PQ15B	E. coli
71 & 72	5'-methyltetrahydropteroyl triglutamate-homocysteine methyl transferase	NMB0944	N. meningitidis
79 & 80	α -1,2 N-acetylglucosamine transferase	RFAK	N. meningitidis
83 & 84	Putative RNA methylase	NMB1348	N. meningitidis
89 & 90	L-lactate permease	NMB0543	N. meningitidis
91 & 92	ABC transporter	NMB1240	N. meningitidis
103 & 104	Probable GTP-binding protein	HI0393	H. influenzae
109 & 110	Capsule polysaccharide modification protein	LIPB	N. meningitidis
113 & 114	Hypothetical 17.8 kD protein	NMB0734	N. meningitidis
115 & 116	<i>E. coli</i> hypothetical protein	YIGC	S. typhimurium
119 & 120	Ribonuclease III	NMB0686	N. meningitidis
121 & 122	AMPD protein	NMB0668	N. meningitidis
123 & 124	5-methyltetrahydropteroyl triglutamate-homocysteine methyl transferase	NMB0944	N. meningitidis

133 & 134	Putative ATP-depenent RNA helicase	NMB1422	N. meningitidis
137 & 138	Putative RNA methylase	NMB1348	N. meningitidis
141 & 142	Shikimate dehydrogenase	AROE	N. meningitidis
143 & 144	Putative outer membrane protein	OMPU	N. meningitidis
145 & 146	TONB protein	TONB	N. meningitidis
147 & 148	Putative apolipoprotein N-acyl transferase	NMB0713	N. meningitidis
149 & 150	Transposase	NMB0991	N. meningitidis
153 & 154	UTP-glucose-1-phosphate uridylyltransferase	NMB0638	N. meningitidis
157 & 158	ADP Heptose-LPS heptosyl transferase II	NMB1527	N. meningitidis
161 & 162	Putative membrane-bound lytic murein transglycosylase B	NMB1279	N. meningitidis
171 & 172	Putative cell-binding factor	NMB0345	N. meningitidis
173 & 174	P-amino benzoate synthetase	PABB	H. pylori J99
177 & 178	5'-methyltetrahydropteroyl triglutamate-homocysteine methyl transferase	NMB0944	N. meningitidis
183 & 184	Conserved hypothetical protein	NMB0183	N. meningitidis
185 & 186	<i>E. coli</i> hypothetical protein	YIGC	S. typhi LT2

For the remaining sequences identified herein, no homology results were obtained.

The gene products were used to produce polyclonal antibodies that were tested in an Elisa assay against various *N. meningitidis* strains to evaluate their effectiveness

5 as a vaccine candidate. The strains used were:

Neisseria meningitidis (B) Type 1000

Neisseria meningitidis (B) Type NGE31

Neisseria meningitidis (B) Type NGH15

Neisseria meningitidis (B) Type SW2 107

10 *Neisseria meningitidis* (B) Type NHG38

Neisseria meningitidis (B) Type NGE28

Neisseria meningitidis (B) Type 2996

This provides information as to the variety of *N. meningitidis* strains that are recognised by these antibodies.

N. Meningitidis was grown on Columbia agar with chocolated horse blood (Oxoid) for 14 hours at 37°C in 5% CO₂. The cells were scraped from agar plates and
5 resuspended in 20ml PBS in a 50ml tube. The cell suspension was heated for 30 minutes at 56°C to kill the bacteria.

A 50 µl sample of the heat-killed *N. Meningitidis* was spread on the Columbia agar with the chocolated horse blood and incubated for 18 hours at 37C, 5%CO₂. This allows confirmation that all *N. Meningitidis* cells have been killed. The OD₆₂₀ of the
10 suspension is adjusted to 0.1 OD units versus PBS.

Elisa with heat killed *N. meningitidis*

Elisa assays were carried out using the heat-killed *N. meningitidis* using the following protocol. Elisa plates were coated overnight with heat-killed cells (50µl of killed bacteria in PBS to each well of 96 well plate and incubated 4°C).

15 Standard Elisa protocols were followed, with all incubations at 37°C for 1 hour. PBS/3% BSA blocking solution, PBS/Tween 0.1% wash solution, anti-rabbit AP conjugate secondary antibody (Sigma) and Sigma Fast P Nitrophenyl phosphate detection reagent (Sigma) were utilised. The data was read at 405nm using an appropriate micro-titre plate reader. The sera used was that available seven days after
20 the first booster vaccination (day 35 after first vaccination).

The antibodies tested were those raised against the gene products identified as SEQ ID NOS. 8, 102, 140, 158 and 202. In each case, the results showed that the anti-sera recognised several different strains of *N. meningitidis* B.

Ex vivo/in vitro screening.

25 Protection against meningococcal disease in humans has been associated with the presence of bacteriocidal antibodies against *N. meningitidis* (Goldscheider *et al.* J. Exp. Med., 1969; 129: 1307-1326). There is also evidence to suggest a correlation between the presence of detectable bacteriocidal activity and protection in an *in vivo* model (reference Martin, J. Bacteriol., 2000; 83: 27-31). Therefore, the antiserum
30 generated was used to evaluate the bacteriocidal activity of the antibodies generated.

The bactericidal assays were performed with pre-immune sera and the corresponding rabbit antiserum raised against the candidate antigens. Commercially available rabbit serum was used as the complement source following pre-screening to eliminate complement only killing. Dulbecco's PBS (Gibco) was used as a buffer where

necessary. The *N. meningitidis* strain MC58, was grown at 37°C (5% CO₂) for 14 hours prior to use in the assay.

200–400 CFU of MC58 in a 50µl volume were incubated in the presence of complement (50µl) with 100µl of serial dilutions of heat-inactivated serum. Samples at
5 time zero were plated to Columbia agar with chocolate horse blood (Oxoid). After incubation for 60 minutes, the number of surviving bacteria was evaluated by plating to Columbia agar with chocolate horse blood (Oxoid). The bacteriocidal activity was expressed in terms of percentage of bacteria surviving after 60 minutes. All samples were tested in duplicate and plated in triplicate. All appropriate positive and negative
10 controls were utilised. In each test sample, the bacteriocidal activity was substantially greater than that for the pre-immune sera.

In vivo screening.

To evaluate the protective efficacy of vaccine candidates, adult mice were immunised with the recombinant proteins identified herein as SEQ ID NOS. 102 and
15 108 and the protective response determined by live bacterial challenge. For each vaccine candidate 15 six week old mice (6 week old balb/C mice) were vaccinated (subcutaneously) with 25µg of antigen on two separate occasions at three week intervals.

One week after the end of the immunisation schedule, the group was challenged
20 with the homologous bacterial strain MC58. The bacteria were inoculated intraperitoneally in a volume of 500µl in Brain Heart Infusion/ 0.5% iron dextran media at a dose of 10⁷ cfu. Previous results have shown that iron is required for initiation of bacteraemic disease in these animals. This model has previously been used to demonstrate the protective efficacy of vaccination (Lissolo *et al.*, Infect. Immun., 1995;
25 63: 884-890).

Control groups included animals vaccinated with adjuvant alone (negative control) or with adjuvant combined with purified PorA (positive control). PorA is an outer membrane protein expressed exclusively by *N. meningitidis* and is the principal target for bactericidal antibodies induced by outer membrane vesicle vaccines. Monoclonal
30 antibodies against PorA have also been shown to passively protect animals in the infant rat model. PorA however varies considerably between strains and so while it elicits some protection when challenged with a homologous strain, it is not an ideal vaccine candidate. Survival was monitored following challenge. The negative control showed no survival after 48 hours. Those vaccinated with PorA showed 6 survivors at 72 hours.

Those vaccinated with the proteins of SEQ ID NOS. 102 and 108 showed 5 and 3 survivors, respectively.

CLAIMS

1. A peptide encoded by an operon including any of the nucleotide sequences identified herein as SEQ ID NOS. 1, 3, 5, 7, 9, 11, 13, 15, 17, 19, 21, 23, 25, 27, 29, 31, 33, 35, 37, 39, 41, 43, 45, 47, 49, 51, 53, 55, 57, 59, 61, 63, 65, 67, 69, 71, 73, 75, 77, 79, 81, 83, 85, 87, 89, 91, 93, 95, 97, 99, 101, 103, 105, 107, 109, 111, 113, 115, 117, 119, 121, 123, 125, 127, 129, 131, 133, 135, 137, 139, 141, 143, 145, 147, 149, 151, 153, 155, 157, 159, 161, 163, 165, 167, 169, 171, 173, 175, 177, 179, 181, 183, 185, 187, 189, 191, 193, 195, 197, 199, 201, 203, 205, 207, of *N. meningitidis*, or a related molecule having at least 40% sequence similarity or identity at the peptide or nucleotide level in a Gram-negative bacterium, or a functional fragment thereof, for therapeutic or diagnostic use.
2. A peptide according to claim 1, wherein the sequence similarity or identity is at least 60%.
3. A peptide according to claim 1 or claim 2, wherein the sequence similarity or identity is at least 90%.
4. A peptide according to claim 1, comprising the amino acid sequence identified herein as SEQ ID NOS. 2, 4, 6, 8, 10, 12, 14, 16, 18, 20, 22, 24, 26, 28, 30, 32, 34, 36, 38, 40, 42, 44, 46, 48, 50, 52, 54, 56, 58, 60, 62, 64, 66, 68, 70, 72, 74, 76, 78, 80, 82, 84, 86, 88, 90, 92, 94, 96, 98, 100, 102, 104, 106, 108, 110, 112, 114, 116, 118, 120, 122, 124, 126, 128, 130, 132, 134, 136, 138, 140, 142, 144, 146, 148, 150, 152, 154, 156, 158, 160, 162, 164, 166, 168, 170, 172, 174, 176, 178, 180, 182, 184, 186, 188, 190, 192, 194, 196, 198, 200, 202, 204, 206 and 208.
5. A polynucleotide encoding a peptide according to any preceding claim, for therapeutic or diagnostic use.
6. A host transformed to express a peptide according to any of claims 1 to 4.
7. An attenuated microorganism comprising a mutation that disrupts the expression of any of the nucleotide sequences defined in claim 1.
8. A microorganism according to claim 7, wherein the mutation is insertional inactivation or a gene deletion.
9. A microorganism according to claim 7 or claim 8, wherein the microorganism is *Neisseria meningitidis*.
10. A microorganism according to any of claims 7 to 9, comprising a second mutation in a second nucleotide sequence.
11. A microorganism according to any of claims 7 to 10, for therapeutic or diagnostic use.

12. A microorganism according to any of claims 7 to 11, comprising a heterologous antigen, therapeutic peptide or nucleic acid.
13. A vaccine comprising a peptide according to any of claims 1 to 4, or the means for its expression.
- 5 14. A vaccine comprising a microorganism according to any of claims 7 to 12.
15. An antibody raised against a peptide according to any of claims 1 to 4.
16. Use of a product according to any of claims 1 to 12, for the manufacture of a medicament for use in the treatment or prevention of a condition associated with infection by *Neisseria* or Gram-negative bacteria.
- 10 17. Use according to claim 16, wherein the condition is meningitis.
18. Use according to claim 16 or claim 17, for veterinary treatment.
19. Use of a product according to any of claims 1 to 12, in a screening assay for the identification of an antimicrobial drug.

SEQUENCE LISTING

<110> Microscience Limited

<120> Virulence Gene and Protein, and Their Use

<130> REP06436WO

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<150> 0011108.8

<151> 2000-05-08

<160> 214

<170> PatentIn Ver. 2.1

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<213> Neisseria meningitidis

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<222> (1)..(1674)

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gaa aag ccg tgg ctg ttg ctg ttg atg gcg ttt gcc tgg ttg tgg ccc   96
Glu Lys Pro Trp Leu Leu Leu Leu Met Ala Phe Ala Trp Leu Trp Pro
          20             25             30

ggc gtg ttt tcc cac gat ttg tgg aat cct gac gaa cct gcc gtc tat   144
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acc gcc gtc gaa gca ctg gca ggc agc ccc acc cct ttg gtt gcc cat   192
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100 105 110	
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130 135 140	
cac ttt ctc aac ccc gct gcc gcc gcc ttt gcc gcc gcc gga ctg gtg	480
His Phe Leu Asn Pro Ala Ala Ala Ala Phe Ala Ala Ala Gly Leu Val	
145 150 155 160	
ctg cac ggt tat tct ttg gct cgc cgg cgc gtg att gcc gcc tct ttt	528
Leu His Gly Tyr Ser Leu Ala Arg Arg Arg Val Ile Ala Ala Ser Phe	
165 170 175	
ctg ctc ggt acg ggt tgg acg ctg atg tcg ttg gca gca gct tat ccg	576
Leu Leu Gly Thr Gly Trp Thr Leu Met Ser Leu Ala Ala Ala Tyr Pro	
180 185 190	
gcg gca ttt gcc ctg atg ctg ccc ctg ccc gtg ctg atg ttt ttc cgt	624
Ala Ala Phe Ala Leu Met Leu Pro Leu Pro Val Leu Met Phe Phe Arg	
195 200 205	
ccg tgg caa agc agg cgt ttg atg ttg acg gca gtc gcc tcg ctt gcc	672
Pro Trp Gln Ser Arg Arg Leu Met Leu Thr Ala Val Ala Ser Leu Ala	
210 215 220	
ttt gcc ctg ccg ctt atg acc gtt tac ccg ctg ctc ttg gca aaa acg	720
Phe Ala Leu Pro Leu Met Thr Val Tyr Pro Leu Leu Leu Ala Lys Thr	
225 230 235 240	
cag ccc gcg ctg ttc gcg caa tgg ctc gac gat cac gtt ttc ggt acg	768
Gln Pro Ala Leu Phe Ala Gln Trp Leu Asp Asp His Val Phe Gly Thr	
245 250 255	
ttc ggc ggc gtg cgg cac att cag acg gca ttc agt ttg ttt tac tat	816
Phe Gly Gly Val Arg His Ile Gln Thr Ala Phe Ser Leu Phe Tyr Tyr	
260 265 270	

ctg aaa aac ctg ctt tgg ttt gca ttg cct gcg ctg ccg ctg gcg gtt	864
Leu Lys Asn Leu Leu Trp Phe Ala Leu Pro Ala Leu Pro Leu Ala Val	
275 280 285	
tgg acg gtt tgc cgc acg cgc ctg ttt tcg acc gac tgg ggg att ttg	912
Trp Thr Val Cys Arg Thr Arg Leu Phe Ser Thr Asp Trp Gly Ile Leu	
290 295 300	
ggc gtc gtc tgg atg ctt gcc gtt ttg gtg ctg ctt gcc gtc aat ccg	960
Gly Val Val Trp Met Leu Ala Val Leu Val Leu Leu Ala Val Asn Pro	
305 310 315 320	
cag cgt ttt cag gat aac ctc gtc tgg ctg ctt ccg ccg ctt gcc ctg	1008
Gln Arg Phe Gln Asp Asn Leu Val Trp Leu Leu Pro Pro Leu Ala Leu	
325 330 335	
ttc ggc gcg gcg caa ctg gac agc ctg aga cgc ggc gcg gcg gcg ttt	1056
Phe Gly Ala Ala Gln Leu Asp Ser Leu Arg Arg Gly Ala Ala Ala Phe	
340 345 350	
gtc aac tgg ttc ggc att atg gcg ttc gga ctg ttt gcc gtg ttc ctg	1104
Val Asn Trp Phe Gly Ile Met Ala Phe Gly Leu Phe Ala Val Phe Leu	
355 360 365	
tgg acg ggc ttt ttc gcc atg aat tac ggc tgg ccc gcc aag ctt gcc	1152
Trp Thr Gly Phe Phe Ala Met Asn Tyr Gly Trp Pro Ala Lys Leu Ala	
370 375 380	
gaa cgc gcc gcc tat ttc agc ccg tat tat gtt cct gat atc gat ccc	1200
Glu Arg Ala Ala Tyr Phe Ser Pro Tyr Tyr Val Pro Asp Ile Asp Pro	
385 390 395 400	
att ccg atg gcg gtt gcc gta ctg ttc aca ccc ttg tgg ctg tgg gcg	1248
Ile Pro Met Ala Val Ala Val Leu Phe Thr Pro Leu Trp Leu Trp Ala	
405 410 415	
att acc cgc aaa aac ata cgc ggc agg cag gcg gtt acc aac tgg gcg	1296
Ile Thr Arg Lys Asn Ile Arg Gly Arg Gln Ala Val Thr Asn Trp Ala	
420 425 430	
gca ggc gtt acc ctg acc tgg gct ttg ctg atg acg ctg ttc ctg ccg	1344
Ala Gly Val Thr Leu Thr Trp Ala Leu Leu Met Thr Leu Phe Leu Pro	
435 440 445	
tgg ctg gac gcg gcg aaa agc cac gcg ccc gtc gtc cgg agt atg gag	1392
Trp Leu Asp Ala Ala Lys Ser His Ala Pro Val Val Arg Ser Met Glu	
450 455 460	

gca tcg ctt tcc ccg gaa tta aaa cgg gag ctt tca gac ggc atc gag 1440
 Ala Ser Leu Ser Pro Glu Leu Lys Arg Glu Leu Ser Asp Gly Ile Glu
 465 470 475 480

tgt atc gac ata ggc ggc ggc gac cta cac acg cgg att gtt tgg acg 1488
 Cys Ile Asp Ile Gly Gly Gly Asp Leu His Thr Arg Ile Val Trp Thr
 485 490 495

cag tac ggc aca ttg ccg cac cgc gtc ggc gat gta caa tgc cgc tac 1536
 Gln Tyr Gly Thr Leu Pro His Arg Val Gly Asp Val Gln Cys Arg Tyr
 500 505 510

cgc atc gtc cgc ttg ccc caa aac gcg gat gcg ccg caa ggc tgg cag 1584
 Arg Ile Val Arg Leu Pro Gln Asn Ala Asp Ala Pro Gln Gly Trp Gln
 515 520 525

acg gtc tgg cag ggt gcg cgc ccg cgc aac aaa gac agt aag ttc gca 1632
 Thr Val Trp Gln Gly Ala Arg Pro Arg Asn Lys Asp Ser Lys Phe Ala
 530 535 540

ctg ata cgg aaa acc ggg gaa aat ata tta aaa aca aca gat 1674
 Leu Ile Arg Lys Thr Gly Glu Asn Ile Leu Lys Thr Thr Asp
 545 550 555

<210> 2

<211> 558

<212> PRT

<213> Neisseria meningitidis

<400> 2

Met Leu Thr Tyr Thr Pro Pro Asp Ala Arg Pro Pro Ala Lys Thr His
 1 5 10 15

Glu Lys Pro Trp Leu Leu Leu Leu Met Ala Phe Ala Trp Leu Trp Pro
 20 25 30

Gly Val Phe Ser His Asp Leu Trp Asn Pro Asp Glu Pro Ala Val Tyr
 35 40 45

Thr Ala Val Glu Ala Leu Ala Gly Ser Pro Thr Pro Leu Val Ala His
 50 55 60

Leu Phe Gly Gln Ile Asp Phe Gly Ile Pro Pro Val Tyr Leu Trp Val
 65 70 75 80

Ala Ala Ala Phe Lys His Leu Leu Ser Pro Trp Ala Ala Asp Pro Tyr
 85 90 95

Asp Ala Ala Arg Phe Ala Gly Val Phe Phe Ala Val Val Gly Leu Thr
 100 105 110

Ser Cys Gly Phe Ala Gly Phe Asn Phe Leu Gly Arg His His Gly Arg
 115 120 125

Ser Val Val Leu Ile Leu Ile Gly Cys Ile Gly Leu Ile Pro Thr Val
 130 135 140

His Phe Leu Asn Pro Ala Ala Ala Phe Ala Ala Ala Gly Leu Val
 145 150 155 160

Leu His Gly Tyr Ser Leu Ala Arg Arg Arg Val Ile Ala Ala Ser Phe
 165 170 175

Leu Leu Gly Thr Gly Trp Thr Leu Met Ser Leu Ala Ala Ala Tyr Pro
 180 185 190

Ala Ala Phe Ala Leu Met Leu Pro Leu Pro Val Leu Met Phe Phe Arg
 195 200 205

Pro Trp Gln Ser Arg Arg Leu Met Leu Thr Ala Val Ala Ser Leu Ala
 210 215 220

Phe Ala Leu Pro Leu Met Thr Val Tyr Pro Leu Leu Leu Ala Lys Thr
 225 230 235 240

Gln Pro Ala Leu Phe Ala Gln Trp Leu Asp Asp His Val Phe Gly Thr
 245 250 255

Phe Gly Gly Val Arg His Ile Gln Thr Ala Phe Ser Leu Phe Tyr Tyr
 260 265 270

Leu Lys Asn Leu Leu Trp Phe Ala Leu Pro Ala Leu Pro Leu Ala Val
 275 280 285

Trp Thr Val Cys Arg Thr Arg Leu Phe Ser Thr Asp Trp Gly Ile Leu
 290 295 300

Gly Val Val Trp Met Leu Ala Val Leu Val Leu Leu Ala Val Asn Pro
 305 310 315 320

Gln Arg Phe Gln Asp Asn Leu Val Trp Leu Leu Pro Pro Leu Ala Leu
 325 330 335

Phe Gly Ala Ala Gln Leu Asp Ser Leu Arg Arg Gly Ala Ala Ala Phe
 340 345 350

Val Asn Trp Phe Gly Ile Met Ala Phe Gly Leu Phe Ala Val Phe Leu
 355 360 365
 Trp Thr Gly Phe Phe Ala Met Asn Tyr Gly Trp Pro Ala Lys Leu Ala
 370 375 380
 Glu Arg Ala Ala Tyr Phe Ser Pro Tyr Tyr Val Pro Asp Ile Asp Pro
 385 390 395 400
 Ile Pro Met Ala Val Ala Val Leu Phe Thr Pro Leu Trp Leu Trp Ala
 405 410 415
 Ile Thr Arg Lys Asn Ile Arg Gly Arg Gln Ala Val Thr Asn Trp Ala
 420 425 430
 Ala Gly Val Thr Leu Thr Trp Ala Leu Leu Met Thr Leu Phe Leu Pro
 435 440 445
 Trp Leu Asp Ala Ala Lys Ser His Ala Pro Val Val Arg Ser Met Glu
 450 455 460
 Ala Ser Leu Ser Pro Glu Leu Lys Arg Glu Leu Ser Asp Gly Ile Glu
 465 470 475 480
 Cys Ile Asp Ile Gly Gly Gly Asp Leu His Thr Arg Ile Val Trp Thr
 485 490 495
 Gln Tyr Gly Thr Leu Pro His Arg Val Gly Asp Val Gln Cys Arg Tyr
 500 505 510
 Arg Ile Val Arg Leu Pro Gln Asn Ala Asp Ala Pro Gln Gly Trp Gln
 515 520 525
 Thr Val Trp Gln Gly Ala Arg Pro Arg Asn Lys Asp Ser Lys Phe Ala
 530 535 540
 Leu Ile Arg Lys Thr Gly Glu Asn Ile Leu Lys Thr Thr Asp
 545 550 555

<210> 3

<211> 1077

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1077)

<400> 3

atg aaa aca ctg acc gta cac acc ccg tcc cac agc tac ccc atc ttt	48
Met Lys Thr Leu Thr Val His Thr Pro Ser His Ser Tyr Pro Ile Phe	
1 5 10 15	
atc ggc aac gga ctg ctg ccg cag gca gga agc ctg ctc aaa ccg cat	96
Ile Gly Asn Gly Leu Leu Pro Gln Ala Gly Ser Leu Leu Lys Pro His	
20 25 30	
ttg ggc aaa cgc gcc gcc atc atc acc aac gaa acc gtc gcc ccg ctc	144
Leu Gly Lys Arg Ala Ala Ile Ile Thr Asn Glu Thr Val Ala Pro Leu	
35 40 45	
tac ctc ggc acg ctt cag acg gca ttg gat gcg gca ggc gta tcc cat	192
Tyr Leu Gly Thr Leu Gln Thr Ala Leu Asp Ala Ala Gly Val Ser His	
50 55 60	
ttc agc atc atc ctg ccc gac ggc gag gcg cac aaa aac tgg cag acg	240
Phe Ser Ile Ile Leu Pro Asp Gly Glu Ala His Lys Asn Trp Gln Thr	
65 70 75 80	
ctc aac ctc atc ttt gac ggg ctg atg caa aac cgc gcc gaa cgc aaa	288
Leu Asn Leu Ile Phe Asp Gly Leu Met Gln Asn Arg Ala Glu Arg Lys	
85 90 95	
acc acg ctc atc gca ctg ggc ggc ggc gtg atc ggc gat atg acg ggt	336
Thr Thr Leu Ile Ala Leu Gly Gly Gly Val Ile Gly Asp Met Thr Gly	
100 105 110	
ttt gcg gcg gct act tat caa cgc ggc gcg cca ttt gtt caa att ccc	384
Phe Ala Ala Ala Thr Tyr Gln Arg Gly Ala Pro Phe Val Gln Ile Pro	
115 120 125	
act act tta tta agt cag gtc gat tcc tcc gtc ggc ggc aaa acc gcc	432
Thr Thr Leu Leu Ser Gln Val Asp Ser Ser Val Gly Gly Lys Thr Ala	
130 135 140	
atc aac cac ccg ctc ggc aaa aat atg atc ggc gcg ttc tac cag ccg	480
Ile Asn His Pro Leu Gly Lys Asn Met Ile Gly Ala Phe Tyr Gln Pro	
145 150 155 160	
cag gcg gtg ctt gcc gac ttg gac acg ctg cac acc ctg ccc gcc cgc	528
Gln Ala Val Leu Ala Asp Leu Asp Thr Leu His Thr Leu Pro Ala Arg	
165 170 175	

gaa ctg tcc gcc ggt atg gcg gaa gtc atc aaa tac ggc acg ctc ggc	576
Glu Leu Ser Ala Gly Met Ala Glu Val Ile Lys Tyr Gly Thr Leu Gly	
180 185 190	
gac att agt ttt ttt gaa tgg ctg gaa cag cat atg ccc gaa ctg atg	624
Asp Ile Ser Phe Phe Glu Trp Leu Glu Gln His Met Pro Glu Leu Met	
195 200 205	
gca ctc gaa cgc gcg ccc ctc atc caa gcc gtg tac cgc tgt tgc caa	672
Ala Leu Glu Arg Ala Pro Leu Ile Gln Ala Val Tyr Arg Cys Cys Gln	
210 215 220	
atg aag gca gac atc gtc gcc caa gac gaa acc gaa cag ggc ata cgc	720
Met Lys Ala Asp Ile Val Ala Gln Asp Glu Thr Glu Gln Gly Ile Arg	
225 230 235 240	
gca tgg ctc aac ctc gga cac acc ttc gga cac gcc att gaa acc gag	768
Ala Trp Leu Asn Leu Gly His Thr Phe Gly His Ala Ile Glu Thr Glu	
245 250 255	
atg ggt tac ggt act tgg ctg cac ggc gag gcc gtc gcc gcc ggc tgc	816
Met Gly Tyr Gly Thr Trp Leu His Gly Glu Ala Val Ala Ala Gly Cys	
260 265 270	
gtg ttg gcg gcg cgt ttg tcc gaa caa ctg ggc aaa atc tcc gcc gca	864
Val Leu Ala Ala Arg Leu Ser Glu Gln Leu Gly Lys Ile Ser Ala Ala	
275 280 285	
gat acc gcg cgg ctc gcc gcc ctg ctc gaa gcc gcc gga ctg ccg tcc	912
Asp Thr Ala Arg Leu Ala Ala Leu Leu Glu Ala Ala Gly Leu Pro Ser	
290 295 300	
gcg cca ccc gtg ttt gcc ttt gaa aaa tgg ctg gaa cac atg agc cac	960
Ala Pro Pro Val Phe Ala Phe Glu Lys Trp Leu Glu His Met Ser His	
305 310 315 320	
gat aaa aaa gtc agc ggc ggc atc atg cgc ttt atc ggt ctg aac cgg	1008
Asp Lys Lys Val Ser Gly Gly Ile Met Arg Phe Ile Gly Leu Asn Arg	
325 330 335	
ctg ggc gaa gcc aac atc acc gaa att acc gac acg gac atc ctc cgc	1056
Leu Gly Glu Ala Asn Ile Thr Glu Ile Thr Asp Thr Asp Ile Leu Arg	
340 345 350	
cgc acc ctg caa ccg tat ctc	1077
Arg Thr Leu Gln Pro Tyr Leu	
355	

<210> 4

<211> 359

<212> PRT

<213> *Neisseria meningitidis*

<400> 4

Met Lys Thr Leu Thr Val His Thr Pro Ser His Ser Tyr Pro Ile Phe
 1 5 10 15

Ile Gly Asn Gly Leu Leu Pro Gln Ala Gly Ser Leu Leu Lys Pro His
 20 25 30

Leu Gly Lys Arg Ala Ala Ile Ile Thr Asn Glu Thr Val Ala Pro Leu
 35 40 45

Tyr Leu Gly Thr Leu Gln Thr Ala Leu Asp Ala Ala Gly Val Ser His
 50 55 60

Phe Ser Ile Ile Leu Pro Asp Gly Glu Ala His Lys Asn Trp Gln Thr
 65 70 75 80

Leu Asn Leu Ile Phe Asp Gly Leu Met Gln Asn Arg Ala Glu Arg Lys
 85 90 95

Thr Thr Leu Ile Ala Leu Gly Gly Gly Val Ile Gly Asp Met Thr Gly
 100 105 110

Phe Ala Ala Ala Thr Tyr Gln Arg Gly Ala Pro Phe Val Gln Ile Pro
 115 120 125

Thr Thr Leu Leu Ser Gln Val Asp Ser Ser Val Gly Gly Lys Thr Ala
 130 135 140

Ile Asn His Pro Leu Gly Lys Asn Met Ile Gly Ala Phe Tyr Gln Pro
 145 150 155 160

Gln Ala Val Leu Ala Asp Leu Asp Thr Leu His Thr Leu Pro Ala Arg
 165 170 175

Glu Leu Ser Ala Gly Met Ala Glu Val Ile Lys Tyr Gly Thr Leu Gly
 180 185 190

Asp Ile Ser Phe Phe Glu Trp Leu Glu Gln His Met Pro Glu Leu Met
 195 200 205

Ala Leu Glu Arg Ala Pro Leu Ile Gln Ala Val Tyr Arg Cys Cys Gln
 210 215 220

Met Lys Ala Asp Ile Val Ala Gln Asp Glu Thr Glu Gln Gly Ile Arg
 225 230 235 240

Ala Trp Leu Asn Leu Gly His Thr Phe Gly His Ala Ile Glu Thr Glu
 245 250 255

Met Gly Tyr Gly Thr Trp Leu His Gly Glu Ala Val Ala Ala Gly Cys
 260 265 270

Val Leu Ala Ala Arg Leu Ser Glu Gln Leu Gly Lys Ile Ser Ala Ala
 275 280 285

Asp Thr Ala Arg Leu Ala Ala Leu Leu Glu Ala Ala Gly Leu Pro Ser
 290 295 300

Ala Pro Pro Val Phe Ala Phe Glu Lys Trp Leu Glu His Met Ser His
 305 310 315 320

Asp Lys Lys Val Ser Gly Gly Ile Met Arg Phe Ile Gly Leu Asn Arg
 325 330 335

Leu Gly Glu Ala Asn Ile Thr Glu Ile Thr Asp Thr Asp Ile Leu Arg
 340 345 350

Arg Thr Leu Gln Pro Tyr Leu
 355

<210> 5

<211> 1053

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1053)

<400> 5

atg aca cac cat tac ccc aca gac gat att aag att aaa gaa gtt aaa 48

Met Thr His His Tyr Pro Thr Asp Asp Ile Lys Ile Lys Glu Val Lys

1 5 10 15

gag ttg ttg ccg ccg att gcc cat ctt tac gag ctg ccg att tcc aaa 96

Glu Leu Leu Pro Pro Ile Ala His Leu Tyr Glu Leu Pro Ile Ser Lys

20 25 30

gag gct tcg ggc ttg gtt cac cgc acc cgt cag gaa att tcc gat ttg	144
Glu Ala Ser Gly Leu Val His Arg Thr Arg Gln Glu Ile Ser Asp Leu	
35 40 45	
gtt cac ggc agg gac aag cgg ctg ttg gtt att atc ggg ccg tgt tcg	192
Val His Gly Arg Asp Lys Arg Leu Leu Val Ile Ile Gly Pro Cys Ser	
50 55 60	
att cac gat ccg aaa gcg gcg ttg gaa tat gcg gag cgt ttg ttg aaa	240
Ile His Asp Pro Lys Ala Ala Leu Glu Tyr Ala Glu Arg Leu Leu Lys	
65 70 75 80	
ctc cgc aag cag tat gaa aac gag ctt ttg att gtg atg cgc gtt tat	288
Leu Arg Lys Gln Tyr Glu Asn Glu Leu Leu Ile Val Met Arg Val Tyr	
85 90 95	
ttc gag aag ccg agg acg acg gtg ggt tgg aaa ggt ttg att aac gac	336
Phe Glu Lys Pro Arg Thr Thr Val Gly Trp Lys Gly Leu Ile Asn Asp	
100 105 110	
ccg cat ttg gac ggt acg ttt gac atc aat ttc ggt ttg cgt cag gca	384
Pro His Leu Asp Gly Thr Phe Asp Ile Asn Phe Gly Leu Arg Gln Ala	
115 120 125	
cgc agc ctg ttg ttg tcg ctg aac aat atg ggt atg cct gcc tct acc	432
Arg Ser Leu Leu Leu Ser Leu Asn Asn Met Gly Met Pro Ala Ser Thr	
130 135 140	
gag ttt ttg gat atg att acg ccg caa tat tat gcg gac ttg att tct	480
Glu Phe Leu Asp Met Ile Thr Pro Gln Tyr Tyr Ala Asp Leu Ile Ser	
145 150 155 160	
tgg ggg gca atc ggc gcg cgg acg acc gaa agc caa gtt cac cgc gaa	528
Trp Gly Ala Ile Gly Ala Arg Thr Thr Glu Ser Gln Val His Arg Glu	
165 170 175	
ttg gca agc ggg ctg tcc tgc ccc gtc ggc ttt aaa aac ggt acg gat	576
Leu Ala Ser Gly Leu Ser Cys Pro Val Gly Phe Lys Asn Gly Thr Asp	
180 185 190	
ggc aat ttg aag att gcc atc gac gca atc ggt gcg gcg agc cat tcg	624
Gly Asn Leu Lys Ile Ala Ile Asp Ala Ile Gly Ala Ala Ser His Ser	
195 200 205	
cat cat ttc ctg tct gta acc aag gcc ggg cat tcc gcc att gtc cat	672
His His Phe Leu Ser Val Thr Lys Ala Gly His Ser Ala Ile Val His	
210 215 220	

acc ggc ggc aat cct gac tgt cat gtc att ttg cgc ggc gga aaa gag 720
 Thr Gly Gly Asn Pro Asp Cys His Val Ile Leu Arg Gly Gly Lys Glu
 225 230 235 240

ccg aat tat gat gcg gga cac gtc agc gag gcg gcg gaa caa ctg cgt 768
 Pro Asn Tyr Asp Ala Gly His Val Ser Glu Ala Ala Glu Gln Leu Arg
 245 250 255

gcg gca ggc gta acc gac aag ctg atg att gat tgc agc cac gcc aac 816
 Ala Ala Gly Val Thr Asp Lys Leu Met Ile Asp Cys Ser His Ala Asn
 260 265 270

agc cgc aag gat tac acc cga cag atg gaa gtg gcg caa gac att gcc 864
 Ser Arg Lys Asp Tyr Thr Arg Gln Met Glu Val Ala Gln Asp Ile Ala
 275 280 285

gcc caa ttg gaa cag gac ggc ggc aat att atg ggc gtg atg gtg gaa 912
 Ala Gln Leu Glu Gln Asp Gly Gly Asn Ile Met Gly Val Met Val Glu
 290 295 300

agc cat ttg gtc gaa ggc agg cag gac aag ccg gaa gtg tac ggc aaa 960
 Ser His Leu Val Glu Gly Arg Gln Asp Lys Pro Glu Val Tyr Gly Lys
 305 310 315 320

agc att acc gat gcg tgt att ggt tgg gac gcg act gaa gaa ctg ttg 1008
 Ser Ile Thr Asp Ala Cys Ile Gly Trp Asp Ala Thr Glu Glu Leu Leu
 325 330 335

gca ttg ttg gca ggt gca aac aaa aaa cgt atg gcg cgc gcc ggt 1053
 Ala Leu Leu Ala Gly Ala Asn Lys Lys Arg Met Ala Arg Ala Gly
 340 345 350

<210> 6

<211> 351

<212> PRT

<213> Neisseria meningitidis

<400> 6

Met Thr His His Tyr Pro Thr Asp Asp Ile Lys Ile Lys Glu Val Lys
 1 5 10 15

Glu Leu Leu Pro Pro Ile Ala His Leu Tyr Glu Leu Pro Ile Ser Lys
 20 25 30

Glu Ala Ser Gly Leu Val His Arg Thr Arg Gln Glu Ile Ser Asp Leu
 35 40 45

Val His Gly Arg Asp Lys Arg Leu Leu Val Ile Ile Gly Pro Cys Ser
 50 55 60

Ile His Asp Pro Lys Ala Ala Leu Glu Tyr Ala Glu Arg Leu Leu Lys
 65 70 75 80

Leu Arg Lys Gln Tyr Glu Asn Glu Leu Leu Ile Val Met Arg Val Tyr
 85 90 95

Phe Glu Lys Pro Arg Thr Thr Val Gly Trp Lys Gly Leu Ile Asn Asp
 100 105 110

Pro His Leu Asp Gly Thr Phe Asp Ile Asn Phe Gly Leu Arg Gln Ala
 115 120 125

Arg Ser Leu Leu Leu Ser Leu Asn Asn Met Gly Met Pro Ala Ser Thr
 130 135 140

Glu Phe Leu Asp Met Ile Thr Pro Gln Tyr Tyr Ala Asp Leu Ile Ser
 145 150 155 160

Trp Gly Ala Ile Gly Ala Arg Thr Thr Glu Ser Gln Val His Arg Glu
 165 170 175

Leu Ala Ser Gly Leu Ser Cys Pro Val Gly Phe Lys Asn Gly Thr Asp
 180 185 190

Gly Asn Leu Lys Ile Ala Ile Asp Ala Ile Gly Ala Ala Ser His Ser
 195 200 205

His His Phe Leu Ser Val Thr Lys Ala Gly His Ser Ala Ile Val His
 210 215 220

Thr Gly Gly Asn Pro Asp Cys His Val Ile Leu Arg Gly Gly Lys Glu
 225 230 235 240

Pro Asn Tyr Asp Ala Gly His Val Ser Glu Ala Ala Glu Gln Leu Arg
 245 250 255

Ala Ala Gly Val Thr Asp Lys Leu Met Ile Asp Cys Ser His Ala Asn
 260 265 270

Ser Arg Lys Asp Tyr Thr Arg Gln Met Glu Val Ala Gln Asp Ile Ala
 275 280 285

Ala Gln Leu Glu Gln Asp Gly Gly Asn Ile Met Gly Val Met Val Glu
 290 295 300

Ser His Leu Val Glu Gly Arg Gln Asp Lys Pro Glu Val Tyr Gly Lys
 305 310 315 320

Ser Ile Thr Asp Ala Cys Ile Gly Trp Asp Ala Thr Glu Glu Leu Leu
 325 330 335

Ala Leu Leu Ala Gly Ala Asn Lys Lys Arg Met Ala Arg Ala Gly
 340 345 350

<210> 7

<211> 1686

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1686)

<400> 7

atg tta caa tcc gaa aat tcg aaa aat tta atc tct tgt tca ata aag 48
 Met Leu Gln Ser Glu Asn Ser Lys Asn Leu Ile Ser Cys Ser Ile Lys
 1 5 10 15

gct tta cca atc atg att tct acc aac ggc atc acc atg cag ttc ggc 96
 Ala Leu Pro Ile Met Ile Ser Thr Asn Gly Ile Thr Met Gln Phe Gly
 20 25 30

gcg aag ccg ctg ttt gaa aac gta tcc gtt aaa ttc ggc gaa ggc aac 144
 Ala Lys Pro Leu Phe Glu Asn Val Ser Val Lys Phe Gly Glu Gly Asn
 35 40 45

cgc tac ggt ttg atc ggc gcg aac ggc tca ggc aaa tcc acc ttc atg 192
 Arg Tyr Gly Leu Ile Gly Ala Asn Gly Ser Gly Lys Ser Thr Phe Met
 50 55 60

aaa atc ctc ggc ggc gat ttg gaa cag aca gcc ggc gaa gtg gcg att 240
 Lys Ile Leu Gly Gly Asp Leu Glu Gln Thr Ala Gly Glu Val Ala Ile
 65 70 75 80

gaa aac ggc gtg cgt ttg ggt aaa ttg cgc caa gac cag ttt gcc tac 288
 Glu Asn Gly Val Arg Leu Gly Lys Leu Arg Gln Asp Gln Phe Ala Tyr
 85 90 95

gaa gat atg cgc gtg ctg gac gtg gta atg atg ggt cac acc gaa atg 336
 Glu Asp Met Arg Val Leu Asp Val Val Met Met Gly His Thr Glu Met
 100 105 110

tgg gcg gcg atg acc gaa cgt gat gcg att tac gcc aat ccc gaa gcc	384
Trp Ala Ala Met Thr Glu Arg Asp Ala Ile Tyr Ala Asn Pro Glu Ala	
115 120 125	
acc gaa gac gac tac atg aaa gcc gca gaa ctg gaa gcc aag ttc gcc	432
Thr Glu Asp Asp Tyr Met Lys Ala Ala Glu Leu Glu Ala Lys Phe Ala	
130 135 140	
gaa tac gac ggc tac acc gcc gaa gcg cgt gcc gcc gag ttg ttg agc	480
Glu Tyr Asp Gly Tyr Thr Ala Glu Ala Arg Ala Ala Glu Leu Leu Ser	
145 150 155 160	
ggc gtg ggc att tcc gaa gat ttg cac aat gca acc atg gca gaa gtt	528
Gly Val Gly Ile Ser Glu Asp Leu His Asn Ala Thr Met Ala Glu Val	
165 170 175	
gcc ccg ggc ttc aaa ctg cgc gta ttg ctg gcg caa gcc ctg ttc tcc	576
Ala Pro Gly Phe Lys Leu Arg Val Leu Leu Ala Gln Ala Leu Phe Ser	
180 185 190	
aag ccg gat gta ttg ctc ttg gac gaa ccg acc aat aac ttg gac att	624
Lys Pro Asp Val Leu Leu Leu Asp Glu Pro Thr Asn Asn Leu Asp Ile	
195 200 205	
aat acc atc cgt tgg ttg gaa ggc gtg ttg aac cag tac gac tcc acg	672
Asn Thr Ile Arg Trp Leu Glu Gly Val Leu Asn Gln Tyr Asp Ser Thr	
210 215 220	
atg att atc atc tcg cac gac cgc cac ttc ttg aac gaa gtc tgc acg	720
Met Ile Ile Ile Ser His Asp Arg His Phe Leu Asn Glu Val Cys Thr	
225 230 235 240	
cat atg gca gat ttg gat tac aac acc atc acc atc tat ccg ggc aac	768
His Met Ala Asp Leu Asp Tyr Asn Thr Ile Thr Ile Tyr Pro Gly Asn	
245 250 255	
tac gac gac tat atg ctc gcc tcc gcc caa tcg cgc gaa cgc gcc ctg	816
Tyr Asp Asp Tyr Met Leu Ala Ser Ala Gln Ser Arg Glu Arg Ala Leu	
260 265 270	
aaa gac aat gcc aag gcg aag gaa aaa ttg cag gaa ttg caa gaa ttt	864
Lys Asp Asn Ala Lys Ala Lys Glu Lys Leu Gln Glu Leu Gln Glu Phe	
275 280 285	
gtg gcg cgg ttc agt gcc aac aaa tcc aaa gcc cgt cag gca acc agc	912
Val Ala Arg Phe Ser Ala Asn Lys Ser Lys Ala Arg Gln Ala Thr Ser	
290 295 300	

cgt	ctg	aaa	cag	gca	gac	aaa	atc	aaa	tcg	gag	atg	gtc	gag	gtc	aaa	960
Arg	Leu	Lys	Gln	Ala	Asp	Lys	Ile	Lys	Ser	Glu	Met	Val	Glu	Val	Lys	
305					310					315					320	
ccc	tct	acc	cgt	caa	aac	ccg	tat	atc	cgt	ttt	gaa	gcc	gat	gaa	aaa	1008
Pro	Ser	Thr	Arg	Gln	Asn	Pro	Tyr	Ile	Arg	Phe	Glu	Ala	Asp	Glu	Lys	
				325					330					335		
gcc	aag	ctg	cac	cgt	cag	gct	gtg	gaa	gtt	gaa	aaa	ctg	gcg	aaa	cgc	1056
Ala	Lys	Leu	His	Arg	Gln	Ala	Val	Glu	Val	Glu	Lys	Leu	Ala	Lys	Arg	
			340					345					350			
ttt	gaa	acc	cag	ttg	ttt	aaa	aac	ctg	aac	ttt	atc	ctt	gaa	gcg	gga	1104
Phe	Glu	Thr	Gln	Leu	Phe	Lys	Asn	Leu	Asn	Phe	Ile	Leu	Glu	Ala	Gly	
		355					360					365				
caa	cgc	ctc	gcc	atc	atc	ggc	ccg	aac	ggc	gcg	ggc	aaa	tcc	acc	ctg	1152
Gln	Arg	Leu	Ala	Ile	Ile	Gly	Pro	Asn	Gly	Ala	Gly	Lys	Ser	Thr	Leu	
	370					375				380						
ctg	aaa	ctc	ttg	gcc	ggc	gcg	tac	aac	ccc	gaa	tat	tca	gac	ggg	ctg	1200
Leu	Lys	Leu	Leu	Ala	Gly	Ala	Tyr	Asn	Pro	Glu	Tyr	Ser	Asp	Gly	Leu	
385					390					395					400	
ctg	ccg	gac	gaa	ggc	agc	atc	aaa	tgg	gcg	gaa	aaa	gcc	agc	gtc	ggc	1248
Leu	Pro	Asp	Glu	Gly	Ser	Ile	Lys	Trp	Ala	Glu	Lys	Ala	Ser	Val	Gly	
			405						410					415		
tac	tat	ccg	caa	gat	cac	gaa	aac	gac	ttc	gat	gtc	gat	atg	gat	ttg	1296
Tyr	Tyr	Pro	Gln	Asp	His	Glu	Asn	Asp	Phe	Asp	Val	Asp	Met	Asp	Leu	
		420						425					430			
agc	gaa	tgg	atg	cgc	caa	tgg	ggg	cag	gac	ggc	gac	gac	gaa	caa	gtc	1344
Ser	Glu	Trp	Met	Arg	Gln	Trp	Gly	Gln	Asp	Gly	Asp	Asp	Glu	Gln	Val	
		435					440					445				
atc	cgc	ggc	act	ttg	ggg	cgt	ttg	ctc	ttc	ggc	agt	aac	gat	gtc	gtg	1392
Ile	Arg	Gly	Thr	Leu	Gly	Arg	Leu	Leu	Phe	Gly	Ser	Asn	Asp	Val	Val	
	450					455					460					
aaa	aaa	gtg	aag	gtt	ctc	tcc	ggg	ggg	gaa	aaa	ggc	cgt	atg	ctt	tac	1440
Lys	Lys	Val	Lys	Val	Leu	Ser	Gly	Gly	Glu	Lys	Gly	Arg	Met	Leu	Tyr	
465					470					475					480	
ggc	aaa	ctg	ttg	ctg	ttg	aaa	ccc	aat	gtc	tta	gtc	atg	gat	gaa	ccg	1488
Gly	Lys	Leu	Leu	Leu	Leu	Lys	Pro	Asn	Val	Leu	Val	Met	Asp	Glu	Pro	
			485						490					495		

acc aac cac atg gac atg gaa agc atc gaa tcc ttg aac atg gcg ctg 1536
 Thr Asn His Met Asp Met Glu Ser Ile Glu Ser Leu Asn Met Ala Leu
 500 505 510

gaa aaa tac aac ggc acg ctg att ttc gtc tcc cac gac cgt cag ttc 1584
 Glu Lys Tyr Asn Gly Thr Leu Ile Phe Val Ser His Asp Arg Gln Phe
 515 520 525

gtt tct tcc ctg gct acc caa atc att gaa ttg gac ggc aaa ggc gga 1632
 Val Ser Ser Leu Ala Thr Gln Ile Ile Glu Leu Asp Gly Lys Gly Gly
 530 535 540

tat gaa cac tac ttg ggc gat tac gaa agc tat ctc gag aaa aaa ggc 1680
 Tyr Glu His Tyr Leu Gly Asp Tyr Glu Ser Tyr Leu Glu Lys Lys Gly
 545 550 555 560

gta gca 1686
 Val Ala

<210> 8

<211> 562

<212> PRT

<213> Neisseria meningitidis

<400> 8

Met Leu Gln Ser Glu Asn Ser Lys Asn Leu Ile Ser Cys Ser Ile Lys
 1 5 10 15

Ala Leu Pro Ile Met Ile Ser Thr Asn Gly Ile Thr Met Gln Phe Gly
 20 25 30

Ala Lys Pro Leu Phe Glu Asn Val Ser Val Lys Phe Gly Glu Gly Asn
 35 40 45

Arg Tyr Gly Leu Ile Gly Ala Asn Gly Ser Gly Lys Ser Thr Phe Met
 50 55 60

Lys Ile Leu Gly Gly Asp Leu Glu Gln Thr Ala Gly Glu Val Ala Ile
 65 70 75 80

Glu Asn Gly Val Arg Leu Gly Lys Leu Arg Gln Asp Gln Phe Ala Tyr
 85 90 95

Glu Asp Met Arg Val Leu Asp Val Val Met Met Gly His Thr Glu Met
 100 105 110

Trp Ala Ala Met Thr Glu Arg Asp Ala Ile Tyr Ala Asn Pro Glu Ala
 115 120 125

Thr Glu Asp Asp Tyr Met Lys Ala Ala Glu Leu Glu Ala Lys Phe Ala
 130 135 140

Glu Tyr Asp Gly Tyr Thr Ala Glu Ala Arg Ala Ala Glu Leu Leu Ser
 145 150 155 160

Gly Val Gly Ile Ser Glu Asp Leu His Asn Ala Thr Met Ala Glu Val
 165 170 175

Ala Pro Gly Phe Lys Leu Arg Val Leu Leu Ala Gln Ala Leu Phe Ser
 180 185 190

Lys Pro Asp Val Leu Leu Leu Asp Glu Pro Thr Asn Asn Leu Asp Ile
 195 200 205

Asn Thr Ile Arg Trp Leu Glu Gly Val Leu Asn Gln Tyr Asp Ser Thr
 210 215 220

Met Ile Ile Ile Ser His Asp Arg His Phe Leu Asn Glu Val Cys Thr
 225 230 235 240

His Met Ala Asp Leu Asp Tyr Asn Thr Ile Thr Ile Tyr Pro Gly Asn
 245 250 255

Tyr Asp Asp Tyr Met Leu Ala Ser Ala Gln Ser Arg Glu Arg Ala Leu
 260 265 270

Lys Asp Asn Ala Lys Ala Lys Glu Lys Leu Gln Glu Leu Gln Glu Phe
 275 280 285

Val Ala Arg Phe Ser Ala Asn Lys Ser Lys Ala Arg Gln Ala Thr Ser
 290 295 300

Arg Leu Lys Gln Ala Asp Lys Ile Lys Ser Glu Met Val Glu Val Lys
 305 310 315 320

Pro Ser Thr Arg Gln Asn Pro Tyr Ile Arg Phe Glu Ala Asp Glu Lys
 325 330 335

Ala Lys Leu His Arg Gln Ala Val Glu Val Glu Lys Leu Ala Lys Arg
 340 345 350

Phe Glu Thr Gln Leu Phe Lys Asn Leu Asn Phe Ile Leu Glu Ala Gly
 355 360 365

Gln Arg Leu Ala Ile Ile Gly Pro Asn Gly Ala Gly Lys Ser Thr Leu
 370 375 380
 Leu Lys Leu Leu Ala Gly Ala Tyr Asn Pro Glu Tyr Ser Asp Gly Leu
 385 390 395 400
 Leu Pro Asp Glu Gly Ser Ile Lys Trp Ala Glu Lys Ala Ser Val Gly
 405 410 415
 Tyr Tyr Pro Gln Asp His Glu Asn Asp Phe Asp Val Asp Met Asp Leu
 420 425 430
 Ser Glu Trp Met Arg Gln Trp Gly Gln Asp Gly Asp Asp Glu Gln Val
 435 440 445
 Ile Arg Gly Thr Leu Gly Arg Leu Leu Phe Gly Ser Asn Asp Val Val
 450 455 460
 Lys Lys Val Lys Val Leu Ser Gly Gly Glu Lys Gly Arg Met Leu Tyr
 465 470 475 480
 Gly Lys Leu Leu Leu Lys Pro Asn Val Leu Val Met Asp Glu Pro
 485 490 495
 Thr Asn His Met Asp Met Glu Ser Ile Glu Ser Leu Asn Met Ala Leu
 500 505 510
 Glu Lys Tyr Asn Gly Thr Leu Ile Phe Val Ser His Asp Arg Gln Phe
 515 520 525
 Val Ser Ser Leu Ala Thr Gln Ile Ile Glu Leu Asp Gly Lys Gly Gly
 530 535 540
 Tyr Glu His Tyr Leu Gly Asp Tyr Glu Ser Tyr Leu Glu Lys Lys Gly
 545 550 555 560
 Val Ala

<210> 9

<211> 1632

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1632)

<400> 9

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Met Ile Lys Pro Asn Leu Arg Pro Lys Leu Gly Ser Ser Ala Leu Ile	
1 5 10 15	
gcc ttc ctt tcc ctg tat tcc tcg ctg gta ttg aat tac gcc ttt ttt	96
Ala Phe Leu Ser Leu Tyr Ser Ser Leu Val Leu Asn Tyr Ala Phe Phe	
20 25 30	
gcc aaa gtt gtc gag ctt cat cct ttt aac ggc acc ggg gcg gat atc	144
Ala Lys Val Val Glu Leu His Pro Phe Asn Gly Thr Gly Ala Asp Ile	
35 40 45	
ttc ctc tat acg atg ccg gtg gtg ctg ttt ttt tta agt aat ttc gtt	192
Phe Leu Tyr Thr Met Pro Val Val Leu Phe Phe Leu Ser Asn Phe Val	
50 55 60	
ttt cac gtc att gcc ctg cct ttc gtg cat aag gta ttg att ccg ttg	240
Phe His Val Ile Ala Leu Pro Phe Val His Lys Val Leu Ile Pro Leu	
65 70 75 80	
ata ttg gtt atc agt gcg gcg gtg tct tac caa gaa ata ttt ttc aat	288
Ile Leu Val Ile Ser Ala Ala Val Ser Tyr Gln Glu Ile Phe Phe Asn	
85 90 95	
atc tat ttc aac aag tcg atg ttg aat aat gtc ttg caa act acg gct	336
Ile Tyr Phe Asn Lys Ser Met Leu Asn Asn Val Leu Gln Thr Thr Ala	
100 105 110	
gcc gaa agc gcg cgc ctg att acg ccg ggc tat gtg ctg tgg att gta	384
Ala Glu Ser Ala Arg Leu Ile Thr Pro Gly Tyr Val Leu Trp Ile Val	
115 120 125	
tgt ttg ggc gta ttg ccc gcg ctg gcg tat atc gcc gtc aag gtt aaa	432
Cys Leu Gly Val Leu Pro Ala Leu Ala Tyr Ile Ala Val Lys Val Lys	
130 135 140	
tac cgc gtt tgg tat aag gag ctt ttg acg cgc ctt gtg ctt gcc gcc	480
Tyr Arg Val Trp Tyr Lys Glu Leu Leu Thr Arg Leu Val Leu Ala Ala	
145 150 155 160	
gtt tcc ttt ttg tgc gcg ttg ggc atc gca atg ttg caa tat cag gat	528
Val Ser Phe Leu Cys Ala Leu Gly Ile Ala Met Leu Gln Tyr Gln Asp	
165 170 175	
tac gcc tcg ttt ttc cgc aac aat aaa tca gta acc cat ctg att gtg	576

Tyr Ala Ser Phe Phe Arg Asn Asn Lys Ser Val Thr His Leu Ile Val	
180 185 190	
cca tct aat ttc atc ggc gcg ggc gtg tcg aaa tac aaa gat tgg aag	624
Pro Ser Asn Phe Ile Gly Ala Gly Val Ser Lys Tyr Lys Asp Trp Lys	
195 200 205	
cgt tcc aat att cct tat acg caa ttg gat atg gcg gtc gtg caa aac	672
Arg Ser Asn Ile Pro Tyr Thr Gln Leu Asp Met Ala Val Val Gln Asn	
210 215 220	
cgc ccg gcc ggc agc ctg cgc cgt ttc gtg gtg ctg gtc gtg ggc gag	720
Arg Pro Ala Gly Ser Leu Arg Arg Phe Val Val Leu Val Val Gly Glu	
225 230 235 240	
acc acg cgt gcc gcc aac tgg ggt ttg aac ggt tac agc cgc caa act	768
Thr Thr Arg Ala Ala Asn Trp Gly Leu Asn Gly Tyr Ser Arg Gln Thr	
245 250 255	
acg ccg ctg ctt gcc gcg cgc ggc gat gaa att gtc aat ttc ccg cag	816
Thr Pro Leu Leu Ala Ala Arg Gly Asp Glu Ile Val Asn Phe Pro Gln	
260 265 270	
gtc aga agc tgc ggt aca tcg acc gcg cac tcc ctg ccg tgt atg ttc	864
Val Arg Ser Cys Gly Thr Ser Thr Ala His Ser Leu Pro Cys Met Phe	
275 280 285	
tca acc ttc gac cgc acg gat tat gac gaa atc aaa gcc gaa cac caa	912
Ser Thr Phe Asp Arg Thr Asp Tyr Asp Glu Ile Lys Ala Glu His Gln	
290 295 300	
gac aac ctg ctg gac atc gtg cag cgc gcc ggc gtg gaa gtt acc tgg	960
Asp Asn Leu Leu Asp Ile Val Gln Arg Ala Gly Val Glu Val Thr Trp	
305 310 315 320	
ttg gaa aac gat tcc ggc tgc aag ggc gtg tgc ggc aaa gtg ccg aat	1008
Leu Glu Asn Asp Ser Gly Cys Lys Gly Val Cys Gly Lys Val Pro Asn	
325 330 335	
acc gac gtt acc tcg ctc aac ctg ccc gaa tac tgc cgc aac ggc gag	1056
Thr Asp Val Thr Ser Leu Asn Leu Pro Glu Tyr Cys Arg Asn Gly Glu	
340 345 350	
tgc ctc gac aat atc ctg ctg acc aag ttc gac gaa gtc ctc aac aaa	1104
Cys Leu Asp Asn Ile Leu Leu Thr Lys Phe Asp Glu Val Leu Asn Lys	
355 360 365	
aac gat aaa gac gcg gtt tta atc ctg cat acc atc ggc agc cac ggg	1152

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Asn Asp Lys Asp Ala Val Leu Ile Leu His Thr Ile Gly Ser His Gly
 370                      375                      380

ccg acg tat tac gaa cgc tat acc gaa gcc gaa cgc aaa ttc acg ccg 1200
Pro Thr Tyr Tyr Glu Arg Tyr Thr Glu Ala Glu Arg Lys Phe Thr Pro
385                      390                      395                      400

acc tgc gac acc aac gaa atc aac aaa tgc acc cgc gcc acg ctg gtc 1248
Thr Cys Asp Thr Asn Glu Ile Asn Lys Cys Thr Arg Ala Thr Leu Val
                      405                      410                      415

aac act tac gac aat acg gtt ttg tat gtg gac cag ttt atc gac aag 1296
Asn Thr Tyr Asp Asn Thr Val Leu Tyr Val Asp Gln Phe Ile Asp Lys
                      420                      425                      430

gtt atc cgc aaa ctt gaa aac cgc gac gat ttg gaa agc gtg gtg cat 1344
Val Ile Arg Lys Leu Glu Asn Arg Asp Asp Leu Glu Ser Val Val His
                      435                      440                      445

tat gtt tcc gac cac ggc gaa agt ttg ggc gaa aac ggg atg tac ctg 1392
Tyr Val Ser Asp His Gly Glu Ser Leu Gly Glu Asn Gly Met Tyr Leu
                      450                      455                      460

cac gcc gcg cct tac gcc atc gcg cct tcc ggg cag acg cat atc ccg 1440
His Ala Ala Pro Tyr Ala Ile Ala Pro Ser Gly Gln Thr His Ile Pro
465                      470                      475                      480

atg gtt atg tgg ttt tcc aaa gcc ttc cgc caa cac ggg ggc ata gat 1488
Met Val Met Trp Phe Ser Lys Ala Phe Arg Gln His Gly Gly Ile Asp
                      485                      490                      495

ttc caa tgc ctc aaa caa aaa gcg gcg gaa aac gaa tat tcg cac gac 1536
Phe Gln Cys Leu Lys Gln Lys Ala Ala Glu Asn Glu Tyr Ser His Asp
                      500                      505                      510

cac tat ttc agc acg gta ttg ggg ctg atg gac att tcc aac agc caa 1584
His Tyr Phe Ser Thr Val Leu Gly Leu Met Asp Ile Ser Asn Ser Gln
                      515                      520                      525

acc tat cgg aag gaa atg gat ata ttg gca gcc tgc cgc cgt ccg cgc 1632
Thr Tyr Arg Lys Glu Met Asp Ile Leu Ala Ala Cys Arg Arg Pro Arg
530                      535                      540

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<210> 10

<211> 544

<212> PRT

<213> Neisseria meningitidis

<400> 10

Met Ile Lys Pro Asn Leu Arg Pro Lys Leu Gly Ser Ser Ala Leu Ile
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Ala Phe Leu Ser Leu Tyr Ser Ser Leu Val Leu Asn Tyr Ala Phe Phe
 20 25 30

Ala Lys Val Val Glu Leu His Pro Phe Asn Gly Thr Gly Ala Asp Ile
 35 40 45

Phe Leu Tyr Thr Met Pro Val Val Leu Phe Phe Leu Ser Asn Phe Val
 50 55 60

Phe His Val Ile Ala Leu Pro Phe Val His Lys Val Leu Ile Pro Leu
 65 70 75 80

Ile Leu Val Ile Ser Ala Ala Val Ser Tyr Gln Glu Ile Phe Phe Asn
 85 90 95

Ile Tyr Phe Asn Lys Ser Met Leu Asn Asn Val Leu Gln Thr Thr Ala
 100 105 110

Ala Glu Ser Ala Arg Leu Ile Thr Pro Gly Tyr Val Leu Trp Ile Val
 115 120 125

Cys Leu Gly Val Leu Pro Ala Leu Ala Tyr Ile Ala Val Lys Val Lys
 130 135 140

Tyr Arg Val Trp Tyr Lys Glu Leu Leu Thr Arg Leu Val Leu Ala Ala
 145 150 155 160

Val Ser Phe Leu Cys Ala Leu Gly Ile Ala Met Leu Gln Tyr Gln Asp
 165 170 175

Tyr Ala Ser Phe Phe Arg Asn Asn Lys Ser Val Thr His Leu Ile Val
 180 185 190

Pro Ser Asn Phe Ile Gly Ala Gly Val Ser Lys Tyr Lys Asp Trp Lys
 195 200 205

Arg Ser Asn Ile Pro Tyr Thr Gln Leu Asp Met Ala Val Val Gln Asn
 210 215 220

Arg Pro Ala Gly Ser Leu Arg Arg Phe Val Val Leu Val Val Gly Glu
 225 230 235 240

Thr Thr Arg Ala Ala Asn Trp Gly Leu Asn Gly Tyr Ser Arg Gln Thr

	245		250		255
Thr Pro Leu Leu Ala Ala Arg Gly Asp Glu Ile Val Asn Phe Pro Gln					
	260		265		270
Val Arg Ser Cys Gly Thr Ser Thr Ala His Ser Leu Pro Cys Met Phe					
	275		280		285
Ser Thr Phe Asp Arg Thr Asp Tyr Asp Glu Ile Lys Ala Glu His Gln					
	290		295		300
Asp Asn Leu Leu Asp Ile Val Gln Arg Ala Gly Val Glu Val Thr Trp					
305		310		315	320
Leu Glu Asn Asp Ser Gly Cys Lys Gly Val Cys Gly Lys Val Pro Asn					
	325		330		335
Thr Asp Val Thr Ser Leu Asn Leu Pro Glu Tyr Cys Arg Asn Gly Glu					
	340		345		350
Cys Leu Asp Asn Ile Leu Leu Thr Lys Phe Asp Glu Val Leu Asn Lys					
	355		360		365
Asn Asp Lys Asp Ala Val Leu Ile Leu His Thr Ile Gly Ser His Gly					
	370		375		380
Pro Thr Tyr Tyr Glu Arg Tyr Thr Glu Ala Glu Arg Lys Phe Thr Pro					
385		390		395	400
Thr Cys Asp Thr Asn Glu Ile Asn Lys Cys Thr Arg Ala Thr Leu Val					
	405		410		415
Asn Thr Tyr Asp Asn Thr Val Leu Tyr Val Asp Gln Phe Ile Asp Lys					
	420		425		430
Val Ile Arg Lys Leu Glu Asn Arg Asp Asp Leu Glu Ser Val Val His					
	435		440		445
Tyr Val Ser Asp His Gly Glu Ser Leu Gly Glu Asn Gly Met Tyr Leu					
	450		455		460
His Ala Ala Pro Tyr Ala Ile Ala Pro Ser Gly Gln Thr His Ile Pro					
465		470		475	480
Met Val Met Trp Phe Ser Lys Ala Phe Arg Gln His Gly Gly Ile Asp					
	485		490		495
Phe Gln Cys Leu Lys Gln Lys Ala Ala Glu Asn Glu Tyr Ser His Asp					

500	505	510
His Tyr Phe Ser Thr Val Leu Gly Leu Met Asp Ile Ser Asn Ser Gln		
515	520	525
Thr Tyr Arg Lys Glu Met Asp Ile Leu Ala Ala Cys Arg Arg Pro Arg		
530	535	540

<210> 11

<211> 1052

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(1050)

<400> 11

atg aca cac cat tac ccc aca gac gat att aag att aaa gaa gtt aaa	48
Met Thr His His Tyr Pro Thr Asp Asp Ile Lys Ile Lys Glu Val Lys	
1 5 10 15	
gag ttg ttg ccg ccg att gcc cat ctt tac gag ctg ccg att tcc aaa	96
Glu Leu Leu Pro Pro Ile Ala His Leu Tyr Glu Leu Pro Ile Ser Lys	
20 25 30	
gag gct tcg ggc ttg gtt cac cgc acc cgt cag gaa att tcc gat ttg	144
Glu Ala Ser Gly Leu Val His Arg Thr Arg Gln Glu Ile Ser Asp Leu	
35 40 45	
gtt cac ggc agg gac aag cgg ctg ttg gtt att atc ggg ccg tgt tcg	192
Val His Gly Arg Asp Lys Arg Leu Leu Val Ile Ile Gly Pro Cys Ser	
50 55 60	
att cac gat ccg aaa gcg gcg ttg gaa tat gcg gag cgt ttg ttg aaa	240
Ile His Asp Pro Lys Ala Ala Leu Glu Tyr Ala Glu Arg Leu Leu Lys	
65 70 75 80	
ctc cgc aag cag tat gaa aac gag ctt ttg att gtg atg cgc gtt tat	288
Leu Arg Lys Gln Tyr Glu Asn Glu Leu Leu Ile Val Met Arg Val Tyr	
85 90 95	
ttc gag aag ccg agg acg acg gtg ggt tgg aaa ggt ttg att aac gac	336
Phe Glu Lys Pro Arg Thr Thr Val Gly Trp Lys Gly Leu Ile Asn Asp	
100 105 110	

ccg cat ttg gac ggt acg ttt gac atc aat ttc ggt ttg cgt cag gca	384
Pro His Leu Asp Gly Thr Phe Asp Ile Asn Phe Gly Leu Arg Gln Ala	
115 120 125	
cgc agc ctg ttg ttg tcg ctg aac aat atg ggt atg cct gcc tct acc	432
Arg Ser Leu Leu Leu Ser Leu Asn Asn Met Gly Met Pro Ala Ser Thr	
130 135 140	
gag ttt ttg gat atg att acg ccg caa tat tat gcg gac ttg att tct	480
Glu Phe Leu Asp Met Ile Thr Pro Gln Tyr Tyr Ala Asp Leu Ile Ser	
145 150 155 160	
tgg ggg gca atc ggc gcg cgg acg acc gaa agc caa gtt cac cgc gaa	528
Trp Gly Ala Ile Gly Ala Arg Thr Thr Glu Ser Gln Val His Arg Glu	
165 170 175	
ttg gca agc ggg ctg tcc tgc ccc gtc ggc ttt aaa aac ggt acg gat	576
Leu Ala Ser Gly Leu Ser Cys Pro Val Gly Phe Lys Asn Gly Thr Asp	
180 185 190	
ggc aat ttg aag att gcc atc gac gca atc ggt gcg gcg agc cat tcg	624
Gly Asn Leu Lys Ile Ala Ile Asp Ala Ile Gly Ala Ala Ser His Ser	
195 200 205	
cat cat ttc ctg tct gta acc aag gcc ggg cat tcc gcc att gtc cat	672
His His Phe Leu Ser Val Thr Lys Ala Gly His Ser Ala Ile Val His	
210 215 220	
acc ggc ggc aat cct gac tgt cat gtc att ttg cgc ggc gga aaa gag	720
Thr Gly Gly Asn Pro Asp Cys His Val Ile Leu Arg Gly Gly Lys Glu	
225 230 235 240	
ccg aat tat gat gcg gga cac gtc agc gag gcg gcg gaa caa ctg cgt	768
Pro Asn Tyr Asp Ala Gly His Val Ser Glu Ala Ala Glu Gln Leu Arg	
245 250 255	
gcg gca ggc gta acc gac aag ctg atg att gat tgc agc cac gcc aac	816
Ala Ala Gly Val Thr Asp Lys Leu Met Ile Asp Cys Ser His Ala Asn	
260 265 270	
agc cgc aag gat tac acc cga cag atg gaa gtg gcg caa gac att gcc	864
Ser Arg Lys Asp Tyr Thr Arg Gln Met Glu Val Ala Gln Asp Ile Ala	
275 280 285	
gcc caa ttg gaa cag gac ggc ggc aat att atg ggc gtg atg gtg gaa	912
Ala Gln Leu Glu Gln Asp Gly Gly Asn Ile Met Gly Val Met Val Glu	
290 295 300	

agc cat ttg gtc gaa ggc agg cag gac aag ccg gaa gtg tac ggc aaa 960
 Ser His Leu Val Glu Gly Arg Gln Asp Lys Pro Glu Val Tyr Gly Lys
 305 310 315 320
 agc att acc gat gcg tgt att ggt tgg gac gcg act gaa gaa ctg ttg 1008
 Ser Ile Thr Asp Ala Cys Ile Gly Trp Asp Ala Thr Glu Glu Leu Leu
 325 330 335
 gca ttg ttg gca ggt gca aac aaa aaa cgt atg gcg cgc gcc gg 1052
 Ala Leu Leu Ala Gly Ala Asn Lys Lys Arg Met Ala Arg Ala
 340 345 350

<210> 12
 <211> 350
 <212> PRT
 <213> Neisseria meningitidis

<400> 12
 Met Thr His His Tyr Pro Thr Asp Asp Ile Lys Ile Lys Glu Val Lys
 1 5 10 15
 Glu Leu Leu Pro Pro Ile Ala His Leu Tyr Glu Leu Pro Ile Ser Lys
 20 25 30
 Glu Ala Ser Gly Leu Val His Arg Thr Arg Gln Glu Ile Ser Asp Leu
 35 40 45
 Val His Gly Arg Asp Lys Arg Leu Leu Val Ile Ile Gly Pro Cys Ser
 50 55 60
 Ile His Asp Pro Lys Ala Ala Leu Glu Tyr Ala Glu Arg Leu Leu Lys
 65 70 75 80
 Leu Arg Lys Gln Tyr Glu Asn Glu Leu Leu Ile Val Met Arg Val Tyr
 85 90 95
 Phe Glu Lys Pro Arg Thr Thr Val Gly Trp Lys Gly Leu Ile Asn Asp
 100 105 110
 Pro His Leu Asp Gly Thr Phe Asp Ile Asn Phe Gly Leu Arg Gln Ala
 115 120 125
 Arg Ser Leu Leu Leu Ser Leu Asn Asn Met Gly Met Pro Ala Ser Thr
 130 135 140
 Glu Phe Leu Asp Met Ile Thr Pro Gln Tyr Tyr Ala Asp Leu Ile Ser
 145 150 155 160

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Trp Gly Ala Ile Gly Ala Arg Thr Thr Glu Ser Gln Val His Arg Glu
      165                      170                      175

Leu Ala Ser Gly Leu Ser Cys Pro Val Gly Phe Lys Asn Gly Thr Asp
      180                      185                      190

Gly Asn Leu Lys Ile Ala Ile Asp Ala Ile Gly Ala Ala Ser His Ser
      195                      200                      205

His His Phe Leu Ser Val Thr Lys Ala Gly His Ser Ala Ile Val His
      210                      215                      220

Thr Gly Gly Asn Pro Asp Cys His Val Ile Leu Arg Gly Gly Lys Glu
      225                      230                      235                      240

Pro Asn Tyr Asp Ala Gly His Val Ser Glu Ala Ala Glu Gln Leu Arg
      245                      250                      255

Ala Ala Gly Val Thr Asp Lys Leu Met Ile Asp Cys Ser His Ala Asn
      260                      265                      270

Ser Arg Lys Asp Tyr Thr Arg Gln Met Glu Val Ala Gln Asp Ile Ala
      275                      280                      285

Ala Gln Leu Glu Gln Asp Gly Gly Asn Ile Met Gly Val Met Val Glu
      290                      295                      300

Ser His Leu Val Glu Gly Arg Gln Asp Lys Pro Glu Val Tyr Gly Lys
      305                      310                      315                      320

Ser Ile Thr Asp Ala Cys Ile Gly Trp Asp Ala Thr Glu Glu Leu Leu
      325                      330                      335

Ala Leu Leu Ala Gly Ala Asn Lys Lys Arg Met Ala Arg Ala
      340                      345                      350

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<210> 13

<211> 432

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(432)

<400> 13

atg gca ttt ggt tgc atg aat tcc ggc gac gat tct ccg atg tcc gac 48
 Met Ala Phe Gly Ser Met Asn Ser Gly Asp Asp Ser Pro Met Ser Asp
 1 5 10 15

atc aac gtt acg ccg ttg gtg gac gtg atg ctg gtg ttg ctg att gtg 96
 Ile Asn Val Thr Pro Leu Val Asp Val Met Leu Val Leu Leu Ile Val
 20 25 30

ttt atg att act atg ccg gtg ctg acg cat tcc atc cct ttg gaa ctg 144
 Phe Met Ile Thr Met Pro Val Leu Thr His Ser Ile Pro Leu Glu Leu
 35 40 45

ccg acc gcg tcc gag cag aca aac aag cag gac aaa cag cct aaa gac 192
 Pro Thr Ala Ser Glu Gln Thr Asn Lys Gln Asp Lys Gln Pro Lys Asp
 50 55 60

ccc ctg cgc ctg acg att gat gcg aac ggc ggc tat tat gtc ggc ggg 240
 Pro Leu Arg Leu Thr Ile Asp Ala Asn Gly Gly Tyr Tyr Val Gly Gly
 65 70 75 80

gat tct gca agc aaa gtg gaa atc ggg gaa gtg gaa agc cgt ctg aaa 288
 Asp Ser Ala Ser Lys Val Glu Ile Gly Glu Val Glu Ser Arg Leu Lys
 85 90 95

gcc gcc aag gag cag aat gaa aac gtg att gtg gcg att gcg gca gac 336
 Ala Ala Lys Glu Gln Asn Glu Asn Val Ile Val Ala Ile Ala Ala Asp
 100 105 110

aag gcg gtg gaa tac gat tat gta aac aaa gct tta gaa gcc gcc cgt 384
 Lys Ala Val Glu Tyr Asp Tyr Val Asn Lys Ala Leu Glu Ala Ala Arg
 115 120 125

cag gca gga atc acc aaa atc ggt ttt gta acc gaa acc aag gcg caa 432
 Gln Ala Gly Ile Thr Lys Ile Gly Phe Val Thr Glu Thr Lys Ala Gln
 130 135 140

<210> 14

<211> 144

<212> PRT

<213> Neisseria meningitidis

<400> 14

Met Ala Phe Gly Ser Met Asn Ser Gly Asp Asp Ser Pro Met Ser Asp
 1 5 10 15

Ile Asn Val Thr Pro Leu Val Asp Val Met Leu Val Leu Leu Ile Val

20 25 30
 Phe Met Ile Thr Met Pro Val Leu Thr His Ser Ile Pro Leu Glu Leu
 35 40 45
 Pro Thr Ala Ser Glu Gln Thr Asn Lys Gln Asp Lys Gln Pro Lys Asp
 50 55 60
 Pro Leu Arg Leu Thr Ile Asp Ala Asn Gly Gly Tyr Tyr Val Gly Gly
 65 70 75 80
 Asp Ser Ala Ser Lys Val Glu Ile Gly Glu Val Glu Ser Arg Leu Lys
 85 90 95
 Ala Ala Lys Glu Gln Asn Glu Asn Val Ile Val Ala Ile Ala Ala Asp
 100 105 110
 Lys Ala Val Glu Tyr Asp Tyr Val Asn Lys Ala Leu Glu Ala Ala Arg
 115 120 125
 Gln Ala Gly Ile Thr Lys Ile Gly Phe Val Thr Glu Thr Lys Ala Gln
 130 135 140

<210> 15

<211> 267

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(267)

<400> 15

atg ctc aaa caa tcc atc gaa atc atc aac aaa ctc gga ctc cac gcc 48
 Met Leu Lys Gln Ser Ile Glu Ile Ile Asn Lys Leu Gly Leu His Ala
 1 5 10 15

cgc gcg tcc aac aag ttc acc caa acc gcg tcc caa ttc aaa agc gaa 96
 Arg Ala Ser Asn Lys Phe Thr Gln Thr Ala Ser Gln Phe Lys Ser Glu
 20 25 30

gtc tgg gtt acg aaa aac gac agc cgc gtc aac ggc aaa agc att atg 144
 Val Trp Val Thr Lys Asn Asp Ser Arg Val Asn Gly Lys Ser Ile Met
 35 40 45

ggg ctg atg atg ctc gcc gcc gcc aag ggt acg gtc atc gaa ctg gag 192

Gly Leu Met Met Leu Ala Ala Ala Lys Gly Thr Val Ile Glu Leu Glu
 50 55 60

acg gac ggc gcg gac gag gcg gaa gcg atg cgc gcc ctg acc gac tta 240
 Thr Asp Gly Ala Asp Glu Ala Glu Ala Met Arg Ala Leu Thr Asp Leu
 65 70 75 80

atc aac ggc tac ttc ggc gag ggc gaa 267
 Ile Asn Gly Tyr Phe Gly Glu Gly Glu
 85

<210> 16
 <211> 89
 <212> PRT
 <213> Neisseria meningitidis

<400> 16
 Met Leu Lys Gln Ser Ile Glu Ile Ile Asn Lys Leu Gly Leu His Ala
 1 5 10 15

Arg Ala Ser Asn Lys Phe Thr Gln Thr Ala Ser Gln Phe Lys Ser Glu
 20 25 30

Val Trp Val Thr Lys Asn Asp Ser Arg Val Asn Gly Lys Ser Ile Met
 35 40 45

Gly Leu Met Met Leu Ala Ala Ala Lys Gly Thr Val Ile Glu Leu Glu
 50 55 60

Thr Asp Gly Ala Asp Glu Ala Glu Ala Met Arg Ala Leu Thr Asp Leu
 65 70 75 80

Ile Asn Gly Tyr Phe Gly Glu Gly Glu
 85

<210> 17
 <211> 633
 <212> DNA
 <213> Neisseria meningitidis

<220>
 <221> CDS
 <222> (1)..(633)

<400> 17

atg ccc atg aac ctt ttt caa aac gcc aaa ttc ttc acg acg gta aac	48
Met Pro Met Asn Leu Phe Gln Asn Ala Lys Phe Phe Thr Thr Val Asn	
1 5 10 15	
cat ctc aaa gac ctg ccg gac aca ccg ctc gaa att gcc ttt gtc ggc	96
His Leu Lys Asp Leu Pro Asp Thr Pro Leu Glu Ile Ala Phe Val Gly	
20 25 30	
agg agc aat gcc gga aaa tcc agt gcc atc aat acc ctg acc aac cat	144
Arg Ser Asn Ala Gly Lys Ser Ser Ala Ile Asn Thr Leu Thr Asn His	
35 40 45	
gtc cgt ctt gcc tac gtt tca aaa aca ccc gga cgg acg cag cat atc	192
Val Arg Leu Ala Tyr Val Ser Lys Thr Pro Gly Arg Thr Gln His Ile	
50 55 60	
aac ttc ttc gag ctg cag aac ggc aat ttt atg gtc gat ttg ccc ggc	240
Asn Phe Phe Glu Leu Gln Asn Gly Asn Phe Met Val Asp Leu Pro Gly	
65 70 75 80	
tac ggt tat gcc caa gtc ccc gaa gca gta cgc gca cat tgg gtc aat	288
Tyr Gly Tyr Ala Gln Val Pro Glu Ala Val Arg Ala His Trp Val Asn	
85 90 95	
ctg ctc ggc gac tat ctg caa cag cgc aaa cag ctt atc ggg ctg gtt	336
Leu Leu Gly Asp Tyr Leu Gln Gln Arg Lys Gln Leu Ile Gly Leu Val	
100 105 110	
ttg att atg gat gcc cgc cat cct tta aaa gaa ctc gac atc cgt atg	384
Leu Ile Met Asp Ala Arg His Pro Leu Lys Glu Leu Asp Ile Arg Met	
115 120 125	
ctg gac ttt ttc cac acg acc ggc aga ccg gtt cac atc ctg ctg tca	432
Leu Asp Phe Phe His Thr Thr Gly Arg Pro Val His Ile Leu Leu Ser	
130 135 140	
aaa gcc gac aaa tta tcc aaa aac gaa cag ata aaa acc ctg tcc caa	480
Lys Ala Asp Lys Leu Ser Lys Asn Glu Gln Ile Lys Thr Leu Ser Gln	
145 150 155 160	
gtc aaa aaa ctg ctc aaa cct tat tcc gac agg caa aac atc agc gta	528
Val Lys Lys Leu Leu Lys Pro Tyr Ser Asp Arg Gln Asn Ile Ser Val	
165 170 175	
cag ctg ttt tcc agc ctg aaa aaa caa ggt att gac gag gcc aac cga	576
Gln Leu Phe Ser Ser Leu Lys Lys Gln Gly Ile Asp Glu Ala Asn Arg	
180 185 190	

act gtc gga agc tgg ttt gac gca gca gat gcc gcc gct tcc tct cca 624
 Thr Val Gly Ser Trp Phe Asp Ala Ala Asp Ala Ala Ala Ser Ser Pro
 195 200 205

aag gaa aac 633
 Lys Glu Asn
 210

<210> 18

<211> 211

<212> PRT

<213> Neisseria meningitidis

<400> 18

Met Pro Met Asn Leu Phe Gln Asn Ala Lys Phe Phe Thr Thr Val Asn
 1 5 10 15

His Leu Lys Asp Leu Pro Asp Thr Pro Leu Glu Ile Ala Phe Val Gly
 20 25 30

Arg Ser Asn Ala Gly Lys Ser Ser Ala Ile Asn Thr Leu Thr Asn His
 35 40 45

Val Arg Leu Ala Tyr Val Ser Lys Thr Pro Gly Arg Thr Gln His Ile
 50 55 60

Asn Phe Phe Glu Leu Gln Asn Gly Asn Phe Met Val Asp Leu Pro Gly
 65 70 75 80

Tyr Gly Tyr Ala Gln Val Pro Glu Ala Val Arg Ala His Trp Val Asn
 85 90 95

Leu Leu Gly Asp Tyr Leu Gln Gln Arg Lys Gln Leu Ile Gly Leu Val
 100 105 110

Leu Ile Met Asp Ala Arg His Pro Leu Lys Glu Leu Asp Ile Arg Met
 115 120 125

Leu Asp Phe Phe His Thr Thr Gly Arg Pro Val His Ile Leu Leu Ser
 130 135 140

Lys Ala Asp Lys Leu Ser Lys Asn Glu Gln Ile Lys Thr Leu Ser Gln
 145 150 155 160

Val Lys Lys Leu Leu Lys Pro Tyr Ser Asp Arg Gln Asn Ile Ser Val
 165 170 175

Gln Leu Phe Ser Ser Leu Lys Lys Gln Gly Ile Asp Glu Ala Asn Arg
 180 185 190

Thr Val Gly Ser Trp Phe Asp Ala Ala Asp Ala Ala Ala Ser Ser Pro
 195 200 205

Lys Glu Asn
 210

<210> 19

<211> 621

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(621)

<400> 19

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 Met Lys Arg Leu Thr Leu Leu Ala Phe Val Leu Ala Ala Gly Ala Val
 1 5 10 15

tcc gca tct ccc aaa gca gac gtg gaa aaa ggc aaa cag gtt gcc gca 96
 Ser Ala Ser Pro Lys Ala Asp Val Glu Lys Gly Lys Gln Val Ala Ala
 20 25 30

acg gtt tgt gcg gct tgc cat gca gca gac ggt aac agc ggc att gcg 144
 Thr Val Cys Ala Ala Cys His Ala Ala Asp Gly Asn Ser Gly Ile Ala
 35 40 45

atg tat ccg cgt ttg gcg gca cag cat act gct tac atc tat cat caa 192
 Met Tyr Pro Arg Leu Ala Ala Gln His Thr Ala Tyr Ile Tyr His Gln
 50 55 60

acc atc ggc atc cgc gac ggt aaa cgc acc cac ggt tcg gca gct gtg 240
 Thr Ile Gly Ile Arg Asp Gly Lys Arg Thr His Gly Ser Ala Ala Val
 65 70 75 80

atg aaa ccg gtg gta atg aat ttg agc gat cag gat att ttg aac gta 288
 Met Lys Pro Val Val Met Asn Leu Ser Asp Gln Asp Ile Leu Asn Val
 85 90 95

tcc gca ttc tat gcc aaa cag cag ccc aaa tcc ggt gaa gcc aat cct 336
 Ser Ala Phe Tyr Ala Lys Gln Gln Pro Lys Ser Gly Glu Ala Asn Pro
 100 105 110

aag gaa aat ccc gaa ttg ggt gcg aaa atc tat cgc ggc ggt ttg agc 384
 Lys Glu Asn Pro Glu Leu Gly Ala Lys Ile Tyr Arg Gly Gly Leu Ser
 115 120 125

gat aaa aaa gtg ccg gcg tgt atg tcc tgc cac ggt ccg agc ggt gcg 432
 Asp Lys Lys Val Pro Ala Cys Met Ser Cys His Gly Pro Ser Gly Ala
 130 135 140

ggt atg ccg ggg ggc gga agc gaa att cag gct tat ccg cgt ttg ggc 480
 Gly Met Pro Gly Gly Gly Ser Glu Ile Gln Ala Tyr Pro Arg Leu Gly
 145 150 155 160

ggt cag cat cag gca tat att gtt gaa cag atg aat gcc tac aag tcc 528
 Gly Gln His Gln Ala Tyr Ile Val Glu Gln Met Asn Ala Tyr Lys Ser
 165 170 175

ggt cag cgt aaa aat acc atc atg gaa gat att gca aac cgt atg tct 576
 Gly Gln Arg Lys Asn Thr Ile Met Glu Asp Ile Ala Asn Arg Met Ser
 180 185 190

gaa gaa gat ttg aaa gcg gtc gcc aac ttt atc caa ggt ttg cgt 621
 Glu Glu Asp Leu Lys Ala Val Ala Asn Phe Ile Gln Gly Leu Arg
 195 200 205

<210> 20

<211> 207

<212> PRT

<213> Neisseria meningitidis

<400> 20

Met Lys Arg Leu Thr Leu Leu Ala Phe Val Leu Ala Ala Gly Ala Val
 1 5 10 15

Ser Ala Ser Pro Lys Ala Asp Val Glu Lys Gly Lys Gln Val Ala Ala
 20 25 30

Thr Val Cys Ala Ala Cys His Ala Ala Asp Gly Asn Ser Gly Ile Ala
 35 40 45

Met Tyr Pro Arg Leu Ala Ala Gln His Thr Ala Tyr Ile Tyr His Gln
 50 55 60

Thr Ile Gly Ile Arg Asp Gly Lys Arg Thr His Gly Ser Ala Ala Val
 65 70 75 80

Met Lys Pro Val Val Met Asn Leu Ser Asp Gln Asp Ile Leu Asn Val

	85		90		95
Ser Ala Phe Tyr Ala Lys Gln Gln Pro Lys Ser Gly Glu Ala Asn Pro					
	100		105		110
Lys Glu Asn Pro Glu Leu Gly Ala Lys Ile Tyr Arg Gly Gly Leu Ser					
	115		120		125
Asp Lys Lys Val Pro Ala Cys Met Ser Cys His Gly Pro Ser Gly Ala					
	130		135		140
Gly Met Pro Gly Gly Gly Ser Glu Ile Gln Ala Tyr Pro Arg Leu Gly					
	145		150		155
					160
Gly Gln His Gln Ala Tyr Ile Val Glu Gln Met Asn Ala Tyr Lys Ser					
	165		170		175
Gly Gln Arg Lys Asn Thr Ile Met Glu Asp Ile Ala Asn Arg Met Ser					
	180		185		190
Glu Glu Asp Leu Lys Ala Val Ala Asn Phe Ile Gln Gly Leu Arg					
	195		200		205

<210> 21
 <211> 765
 <212> DNA
 <213> Neisseria meningitidis

<220>
 <221> CDS
 <222> (1)..(765)

<400> 21
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 Met Pro Ser Glu Ala Arg Phe Phe Arg Leu Gln Thr Ala Phe Pro Arg
 1 5 10 15

ttg att tgc ggt ata atc cgc ctt tac cca ttg ttt gca aag cac aat 96
 Leu Ile Cys Gly Ile Ile Arg Leu Tyr Pro Leu Phe Ala Lys His Asn
 20 25 30

atg aca cgc aaa atc ctc gtt acc tcc gcc ctg ccc tat gcc aac ggc 144
 Met Thr Arg Lys Ile Leu Val Thr Ser Ala Leu Pro Tyr Ala Asn Gly
 35 40 45

agc atc cac ctc ggc cac atg gtc gaa cac atc caa acc gac gtt tgg 192

Ser	Ile	His	Leu	Gly	His	Met	Val	Glu	His	Ile	Gln	Thr	Asp	Val	Trp		
50						55					60						
gtg	cgc	ttt	caa	aaa	ctg	cgc	ggc	aac	gaa	tgc	cac	tac	tgc	tgc	gcc	240	
Val	Arg	Phe	Gln	Lys	Leu	Arg	Gly	Asn	Glu	Cys	His	Tyr	Cys	Cys	Ala		
65					70				75						80		
gac	gac	acc	cac	ggc	acg	ccc	gtg	atg	ctt	gcc	gcg	caa	aaa	caa	ggc	288	
Asp	Asp	Thr	His	Gly	Thr	Pro	Val	Met	Leu	Ala	Ala	Gln	Lys	Gln	Gly		
			85					90						95			
atc	gcg	ccc	gaa	gac	atg	att	gcc	aaa	gtg	cgc	gaa	gag	cac	ctc	gcc	336	
Ile	Ala	Pro	Glu	Asp	Met	Ile	Ala	Lys	Val	Arg	Glu	Glu	His	Leu	Ala		
			100					105						110			
gac	ttt	act	ggc	ttt	ttc	atc	ggt	tac	gac	aat	tat	tac	agc	acc	cat	384	
Asp	Phe	Thr	Gly	Phe	Phe	Ile	Gly	Tyr	Asp	Asn	Tyr	Tyr	Ser	Thr	His		
		115					120					125					
tcc	cct	gaa	aac	aaa	cag	ttt	tcc	caa	gac	att	tac	cgc	gcg	ctg	aaa	432	
Ser	Pro	Glu	Asn	Lys	Gln	Phe	Ser	Gln	Asp	Ile	Tyr	Arg	Ala	Leu	Lys		
		130				135					140						
gcc	aac	ggc	aag	att	gag	agc	cgc	gtc	atc	gag	cag	ctt	ttc	gac	ccc	480	
Ala	Asn	Gly	Lys	Ile	Glu	Ser	Arg	Val	Ile	Glu	Gln	Leu	Phe	Asp	Pro		
145					150					155				160			
gaa	aaa	caa	atg	ttc	ctg	ccc	gac	cgc	ttc	gtt	aaa	ggc	gaa	tgc	ccg	528	
Glu	Lys	Gln	Met	Phe	Leu	Pro	Asp	Arg	Phe	Val	Lys	Gly	Glu	Cys	Pro		
			165					170						175			
aaa	tgc	cac	gcc	caa	gac	caa	tac	ggc	gac	aac	tgc	gaa	gtc	tgc	ggc	576	
Lys	Cys	His	Ala	Gln	Asp	Gln	Tyr	Gly	Asp	Asn	Cys	Glu	Val	Cys	Gly		
			180					185						190			
acg	acc	tat	tcc	ccg	acc	gaa	ctg	att	aac	ccg	tat	tcc	gcc	gtt	tcc	624	
Thr	Thr	Tyr	Ser	Pro	Thr	Glu	Leu	Ile	Asn	Pro	Tyr	Ser	Ala	Val	Ser		
		195					200					205					
ggc	gcg	aaa	ccc	gaa	ttg	cgc	gaa	tcc	gaa	cac	ttc	ttc	ttc	aaa	ctg	672	
Gly	Ala	Lys	Pro	Glu	Leu	Arg	Glu	Ser	Glu	His	Phe	Phe	Phe	Lys	Leu		
		210				215					220						
gga	gaa	tgc	gcc	gac	ttc	ctc	aaa	gca	tgg	act	tcc	ggc	aac	aac	ccg	720	
Gly	Glu	Cys	Ala	Asp	Phe	Leu	Lys	Ala	Trp	Thr	Ser	Gly	Asn	Asn	Pro		
225					230					235				240			
cac	gac	ggc	aag	ccc	cat	ctg	caa	gcc	gaa	gcc	ctc	aac	aaa	atg		765	

His Asp Gly Lys Pro His Leu Gln Ala Glu Ala Leu Asn Lys Met
 245 250 255

<210> 22

<211> 255

<212> PRT

<213> Neisseria meningitidis

<400> 22

Met Pro Ser Glu Ala Arg Phe Phe Arg Leu Gln Thr Ala Phe Pro Arg
 1 5 10 15

Leu Ile Cys Gly Ile Ile Arg Leu Tyr Pro Leu Phe Ala Lys His Asn
 20 25 30

Met Thr Arg Lys Ile Leu Val Thr Ser Ala Leu Pro Tyr Ala Asn Gly
 35 40 45

Ser Ile His Leu Gly His Met Val Glu His Ile Gln Thr Asp Val Trp
 50 55 60

Val Arg Phe Gln Lys Leu Arg Gly Asn Glu Cys His Tyr Cys Cys Ala
 65 70 75 80

Asp Asp Thr His Gly Thr Pro Val Met Leu Ala Ala Gln Lys Gln Gly
 85 90 95

Ile Ala Pro Glu Asp Met Ile Ala Lys Val Arg Glu Glu His Leu Ala
 100 105 110

Asp Phe Thr Gly Phe Phe Ile Gly Tyr Asp Asn Tyr Tyr Ser Thr His
 115 120 125

Ser Pro Glu Asn Lys Gln Phe Ser Gln Asp Ile Tyr Arg Ala Leu Lys
 130 135 140

Ala Asn Gly Lys Ile Glu Ser Arg Val Ile Glu Gln Leu Phe Asp Pro
 145 150 155 160

Glu Lys Gln Met Phe Leu Pro Asp Arg Phe Val Lys Gly Glu Cys Pro
 165 170 175

Lys Cys His Ala Gln Asp Gln Tyr Gly Asp Asn Cys Glu Val Cys Gly
 180 185 190

Thr Thr Tyr Ser Pro Thr Glu Leu Ile Asn Pro Tyr Ser Ala Val Ser
 195 200 205

Gly Ala Lys Pro Glu Leu Arg Glu Ser Glu His Phe Phe Phe Lys Leu
 210 215 220

Gly Glu Cys Ala Asp Phe Leu Lys Ala Trp Thr Ser Gly Asn Asn Pro
 225 230 235 240

His Asp Gly Lys Pro His Leu Gln Ala Glu Ala Leu Asn Lys Met
 245 250 255

<210> 23

<211> 1539

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1539)

<400> 23

atg aag ttt tta gac cgt gag gca agc att gcc aag ccg ggt ttc aac 48
 Met Lys Phe Leu Asp Arg Glu Ala Ser Ile Ala Lys Pro Gly Phe Asn
 1 5 10 15

cgt tgg ctc gta ccg ccc gca gca ttg gcg gta cac ctt gcc atc ggg 96
 Arg Trp Leu Val Pro Pro Ala Ala Leu Ala Val His Leu Ala Ile Gly
 20 25 30

cag att tac gcc tat tcc gta ttc aac gcg ccg ctg acc aaa ctt atc 144
 Gln Ile Tyr Ala Tyr Ser Val Phe Asn Ala Pro Leu Thr Lys Leu Ile
 35 40 45

ggc ata acc gaa tcg gcg gcc gga gat tgg aag ctg acg acc gta ggt 192
 Gly Ile Thr Glu Ser Ala Ala Gly Asp Trp Lys Leu Thr Thr Val Gly
 50 55 60

tgg att ttc agt atc gca ctc gcg atg ctg ggc gcg tcg gcc gcc ctg 240
 Trp Ile Phe Ser Ile Ala Leu Ala Met Leu Gly Ala Ser Ala Ala Leu
 65 70 75 80

ttc ggt acg tgg atg gaa cgg gta ggg ccg cgc aaa gcc ata ttt gcc 288
 Phe Gly Thr Trp Met Glu Arg Val Gly Pro Arg Lys Ala Ile Phe Ala
 85 90 95

gcc gcc tgc tgt ttc agc ctg ggc ttc ttc gta tcc gca ttc ggc gtg 336
 Ala Ala Cys Cys Phe Ser Leu Gly Phe Phe Val Ser Ala Phe Gly Val

100	105	110	
cgt acg cac aac ctc ttc ctg ctc tat ttg ggc aac ggc gtc atc ggc			384
Arg Thr His Asn Leu Phe Leu Leu Tyr Leu Gly Asn Gly Val Ile Gly			
115	120	125	
ggc gta ggc ttg ggt ttg ggc tac atc ggg ccg gtt tcc aca ctg atg			432
Gly Val Gly Leu Gly Leu Gly Tyr Ile Gly Pro Val Ser Thr Leu Met			
130	135	140	
aaa tgg ttt ccc gac aag ccg ggt atg gcg acc ggt ttg gcg att atg			480
Lys Trp Phe Pro Asp Lys Pro Gly Met Ala Thr Gly Leu Ala Ile Met			
145	150	155	160
ggg ttc ggt ggc ggc gca atg ctg gcc tcg ccg ctt tcc gta tca ctg			528
Gly Phe Gly Gly Gly Ala Met Leu Ala Ser Pro Leu Ser Val Ser Leu			
165	170	175	
atg aac acc ttt tca aac gct act tcg gtt ggg gtt gcc gaa acc ttc			576
Met Asn Thr Phe Ser Asn Ala Thr Ser Val Gly Val Ala Glu Thr Phe			
180	185	190	
gcc gta ttg ggt ctg ctt tac ctc gca ctg atg atg ttc ggc gcg ttt			624
Ala Val Leu Gly Leu Leu Tyr Leu Ala Leu Met Met Phe Gly Ala Phe			
195	200	205	
acc atc cgc gta cct gcc gac ggc tgg aaa cct gaa ggc tat acc gcg			672
Thr Ile Arg Val Pro Ala Asp Gly Trp Lys Pro Glu Gly Tyr Thr Ala			
210	215	220	
ccg aaa acg caa aac aag ctg gtc agc agc aat cat gtc aac gtg tcc			720
Pro Lys Thr Gln Asn Lys Leu Val Ser Ser Asn His Val Asn Val Ser			
225	230	235	240
caa gcc atg aaa acg ccg cag ttt tgg ctg ttg ttc tgg gta ttg tgc			768
Gln Ala Met Lys Thr Pro Gln Phe Trp Leu Leu Phe Trp Val Leu Cys			
245	250	255	
ctg aac gta act gcc ggc atc ggc gta ttg ggt cag gca tcc gtg atg			816
Leu Asn Val Thr Ala Gly Ile Gly Val Leu Gly Gln Ala Ser Val Met			
260	265	270	
att cag gaa ctc ttt tcc gaa acc tcc ata ggc agg cag gcg gca gtc			864
Ile Gln Glu Leu Phe Ser Glu Thr Ser Ile Gly Arg Gln Ala Ala Val			
275	280	285	
ggc gca ggt gcg gcg gca ggc ttc gtc agc ctc ctg agc ctg ttt aac			912
Gly Ala Gly Ala Ala Ala Gly Phe Val Ser Leu Leu Ser Leu Phe Asn			

290	295	300	
atg ggc gga cgc ttt ttg tgg tgc agc gtt tcc gac aaa atc gga cgc			960
Met Gly Gly Arg Phe Leu Trp Ser Ser Val Ser Asp Lys Ile Gly Arg			
305	310	315	320
aaa aac acc tac acc atc ttc ttc gta ctc ggt tgc ctg ctg tat ttc			1008
Lys Asn Thr Tyr Thr Ile Phe Phe Val Leu Gly Ser Leu Leu Tyr Phe			
	325	330	335
gcc gtt ccc tcc atc ggc gag ggc ggc agc aaa gcc ctg ttc atc atc			1056
Ala Val Pro Ser Ile Gly Glu Gly Gly Ser Lys Ala Leu Phe Ile Ile			
	340	345	350
ggc ttt tgc gtg att att tcc atg tac ggc ggc gga ttt gcc gcc att			1104
Gly Phe Cys Val Ile Ile Ser Met Tyr Gly Gly Gly Phe Ala Ala Ile			
	355	360	365
ccc gcc tac ctg aaa gac ctg ttc ggc acc tat cag gtc ggc gcg att			1152
Pro Ala Tyr Leu Lys Asp Leu Phe Gly Thr Tyr Gln Val Gly Ala Ile			
	370	375	380
cac gga cgc atc ctg ctg gcc tgg tca acc gcc gcc gtc atc ggc ccg			1200
His Gly Arg Ile Leu Leu Ala Trp Ser Thr Ala Ala Val Ile Gly Pro			
385	390	395	400
gta ctg gtc aat tac ata cgc caa agc caa atc gac agc ggc ata ccg			1248
Val Leu Val Asn Tyr Ile Arg Gln Ser Gln Ile Asp Ser Gly Ile Pro			
	405	410	415
gcg gca cag gca tac agc gta acc atg tac atc atg gcg ggt ctg ctg			1296
Ala Ala Gln Ala Tyr Ser Val Thr Met Tyr Ile Met Ala Gly Leu Leu			
	420	425	430
att atc ggt ttg ttg tgc aac ctc gcc gtc aaa tcc gta cac gaa aaa			1344
Ile Ile Gly Leu Leu Cys Asn Leu Ala Val Lys Ser Val His Glu Lys			
	435	440	445
cac cac gaa aaa gac atc aaa acc gcc gca cac agc ggc aac ccc gat			1392
His His Glu Lys Asp Ile Lys Thr Ala Ala His Ser Gly Asn Pro Asp			
	450	455	460
gac gaa acc gcc ata tcc gac gct tac ctt gtg ggc gaa aaa gtt tcc			1440
Asp Glu Thr Ala Ile Ser Asp Ala Tyr Leu Val Gly Glu Lys Val Ser			
465	470	475	480
ggc ggc ggc ata tgc gtc tgg tgg cgt tgg gca ttg gcc gtc ata ccg			1488
Gly Gly Gly Ile Ser Val Trp Trp Arg Trp Ala Leu Ala Val Ile Pro			

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<210> 24
<211> 513
<212> PRT
<213> Neisseria meningitidis
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42

Met Asn Thr Phe Ser Asn Ala Thr Ser Val Gly Val Ala Glu Thr Phe
180 185 190

Ala Val Leu Gly Leu Leu Tyr Leu Ala Leu Met Met Phe Gly Ala Phe
195 200 205

Thr Ile Arg Val Pro Ala Asp Gly Trp Lys Pro Glu Gly Tyr Thr Ala
210 215 220

Pro Lys Thr Gln Asn Lys Leu Val Ser Ser Asn His Val Asn Val Ser
225 230 235 240

Gln Ala Met Lys Thr Pro Gln Phe Trp Leu Leu Phe Trp Val Leu Cys
245 250 255

Leu Asn Val Thr Ala Gly Ile Gly Val Leu Gly Gln Ala Ser Val Met
260 265 270

Ile Gln Glu Leu Phe Ser Glu Thr Ser Ile Gly Arg Gln Ala Ala Val
275 280 285

Gly Ala Gly Ala Ala Ala Gly Phe Val Ser Leu Leu Ser Leu Phe Asn
290 295 300

Met Gly Gly Arg Phe Leu Trp Ser Ser Val Ser Asp Lys Ile Gly Arg
305 310 315 320

Lys Asn Thr Tyr Thr Ile Phe Phe Val Leu Gly Ser Leu Leu Tyr Phe
325 330 335

Ala Val Pro Ser Ile Gly Glu Gly Gly Ser Lys Ala Leu Phe Ile Ile
340 345 350

Gly Phe Cys Val Ile Ile Ser Met Tyr Gly Gly Gly Phe Ala Ala Ile
355 360 365

Pro Ala Tyr Leu Lys Asp Leu Phe Gly Thr Tyr Gln Val Gly Ala Ile
370 375 380

His Gly Arg Ile Leu Leu Ala Trp Ser Thr Ala Ala Val Ile Gly Pro
385 390 395 400

Val Leu Val Asn Tyr Ile Arg Gln Ser Gln Ile Asp Ser Gly Ile Pro
405 410 415

Ala Ala Gln Ala Tyr Ser Val Thr Met Tyr Ile Met Ala Gly Leu Leu
420 425 430

Ile Ile Gly Leu Leu Cys Asn Leu Ala Val Lys Ser Val His Glu Lys
 435 440 445

His His Glu Lys Asp Ile Lys Thr Ala Ala His Ser Gly Asn Pro Asp
 450 455 460

Asp Glu Thr Ala Ile Ser Asp Ala Tyr Leu Val Gly Glu Lys Val Ser
 465 470 475 480

Gly Gly Gly Ile Ser Val Trp Trp Arg Trp Ala Leu Ala Val Ile Pro
 485 490 495

Leu Ala Tyr Gly Val Val Met Ile Phe Val Lys Ala Leu Asp Leu Phe
 500 505 510

Ser

<210> 25

<211> 1038

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1038)

<400> 25

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 Leu Gln Pro Leu Val Ser Val Leu Ile Cys Ala Tyr Asn Val Glu Lys
 1 5 10 15

tat ttc gcc caa tca tta gcc acc gtc gtg aat cag act tgg cgc aac 96
 Tyr Phe Ala Gln Ser Leu Ala Thr Val Val Asn Gln Thr Trp Arg Asn
 20 25 30

ttg gag att ttg att gtc gat gac ggc tcg aca gac ggt acg ctt gcc 144
 Leu Glu Ile Leu Ile Val Asp Asp Gly Ser Thr Asp Gly Thr Leu Ala
 35 40 45

att gcc aag gat ttt caa aag cgg gac agc cgt atc aaa atc ctt gca 192
 Ile Ala Lys Asp Phe Gln Lys Arg Asp Ser Arg Ile Lys Ile Leu Ala
 50 55 60

caa gct caa aat tcc ggc ctg att ccc tct tta aac atc ggg ctg gac 240

45

Asp Ser Leu Glu Tyr Arg Gln Ile Lys Ala Val Ala Tyr Glu Leu Leu
 260 265 270

gag aaa cat ttg ccg gaa gaa gat ttt gaa cgc gcc cgc cgg ttt ttg 864
 Glu Lys His Leu Pro Glu Glu Asp Phe Glu Arg Ala Arg Arg Phe Leu
 275 280 285

tac caa tgc ttc aaa cgg acg gac acg ccg ccc gcc ggc gcg tgg ctg 912
 Tyr Gln Cys Phe Lys Arg Thr Asp Thr Pro Pro Ala Gly Ala Trp Leu
 290 295 300

gat ttc gcg gca gac ggc aag atg agg cgg ctg ttt acc atg agg caa 960
 Asp Phe Ala Ala Asp Gly Lys Met Arg Arg Leu Phe Thr Met Arg Gln
 305 310 315 320

tac ttc ggc att ttg cac cgg ctg att aaa aac cgc cgg cag gcg cgg 1008
 Tyr Phe Gly Ile Leu His Arg Leu Ile Lys Asn Arg Arg Gln Ala Arg
 325 330 335

tcg gat tcg gca ggg aaa gaa cag gag att 1038
 Ser Asp Ser Ala Gly Lys Glu Gln Glu Ile
 340 345

<210> 26
 <211> 346
 <212> PRT
 <213> Neisseria meningitidis

<400> 26
 Leu Gln Pro Leu Val Ser Val Leu Ile Cys Ala Tyr Asn Val Glu Lys
 1 5 10 15

Tyr Phe Ala Gln Ser Leu Ala Thr Val Val Asn Gln Thr Trp Arg Asn
 20 25 30

Leu Glu Ile Leu Ile Val Asp Asp Gly Ser Thr Asp Gly Thr Leu Ala
 35 40 45

Ile Ala Lys Asp Phe Gln Lys Arg Asp Ser Arg Ile Lys Ile Leu Ala
 50 55 60

Gln Ala Gln Asn Ser Gly Leu Ile Pro Ser Leu Asn Ile Gly Leu Asp
 65 70 75 80

Glu Leu Ala Lys Ser Gly Met Gly Glu Tyr Ile Ala Arg Thr Asp Ala
 85 90 95

Asp Asp Ile Ala Ala Pro Asp Trp Ile Glu Lys Ile Val Gly Glu Met
 100 105 110
 Glu Lys Asp Arg Ser Ile Ile Ala Met Gly Ala Trp Leu Glu Val Leu
 115 120 125
 Ser Glu Glu Lys Asp Gly Asn Arg Leu Ala Arg His His Arg His Gly
 130 135 140
 Lys Ile Trp Lys Lys Pro Thr Arg His Glu Asp Ile Ala Asp Phe Phe
 145 150 155 160
 Pro Phe Gly Asn Pro Ile His Asn Asn Thr Met Ile Met Arg Arg Ser
 165 170 175
 Val Ile Asp Gly Gly Leu Arg Tyr Asn Thr Glu Arg Asp Trp Ala Glu
 180 185 190
 Asp Tyr Gln Phe Trp Tyr Asp Val Ser Lys Leu Gly Arg Leu Ala Tyr
 195 200 205
 Tyr Pro Glu Ala Leu Val Lys Tyr Arg Leu His Ala Asn Gln Val Ser
 210 215 220
 Ser Lys Tyr Ser Ile Arg Gln His Glu Ile Ala Gln Gly Ile Gln Lys
 225 230 235 240
 Thr Ala Arg Asn Asp Phe Leu Gln Ser Met Gly Phe Lys Thr Arg Phe
 245 250 255
 Asp Ser Leu Glu Tyr Arg Gln Ile Lys Ala Val Ala Tyr Glu Leu Leu
 260 265 270
 Glu Lys His Leu Pro Glu Glu Asp Phe Glu Arg Ala Arg Arg Phe Leu
 275 280 285
 Tyr Gln Cys Phe Lys Arg Thr Asp Thr Pro Pro Ala Gly Ala Trp Leu
 290 295 300
 Asp Phe Ala Ala Asp Gly Lys Met Arg Arg Leu Phe Thr Met Arg Gln
 305 310 315 320
 Tyr Phe Gly Ile Leu His Arg Leu Ile Lys Asn Arg Arg Gln Ala Arg
 325 330 335
 Ser Asp Ser Ala Gly Lys Glu Gln Glu Ile
 340 345

<210> 27

<211> 1856

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(1857)

<400> 27

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ggc gcg cgc gcg ttg tgg cgc gcc acc ggc gtg atg gaa acc gac ttc      96
Gly Ala Arg Ala Leu Trp Arg Ala Thr Gly Val Met Glu Thr Asp Phe
          20             25             30

ggc aaa ccg att atc gcc gtt gcc aac tcg ttc acc caa ttc gtg cca      144
Gly Lys Pro Ile Ile Ala Val Ala Asn Ser Phe Thr Gln Phe Val Pro
          35             40             45

ggc cat gtc cac ctg cac aat atg ggc cag ctg gtt gcc cgc gaa atc      192
Gly His Val His Leu His Asn Met Gly Gln Leu Val Ala Arg Glu Ile
          50             55             60

gaa aaa gcc ggc gcc atc gcc aaa gaa ttc aac acc atc gcc atc gac      240
Glu Lys Ala Gly Ala Ile Ala Lys Glu Phe Asn Thr Ile Ala Ile Asp
          65             70             75             80

gac ggc atc gct atg gga cac agc ggt atg ctg tat tcc ctg ccc agc      288
Asp Gly Ile Ala Met Gly His Ser Gly Met Leu Tyr Ser Leu Pro Ser
          85             90             95

cgc gat ttg att gcc gac tcc atc gaa tat atg gtc aat gcc cac tgc      336
Arg Asp Leu Ile Ala Asp Ser Ile Glu Tyr Met Val Asn Ala His Cys
          100            105            110

gcc gac gcg ctg gtg tgc att tcc aac tgc gac aaa atc acc ccg ggc      384
Ala Asp Ala Leu Val Cys Ile Ser Asn Cys Asp Lys Ile Thr Pro Gly
          115            120            125

atg ctg att gcc gcg atg cgc ctg aac atc ccg acc atc ttc gtt tcc      432
Met Leu Ile Ala Ala Met Arg Leu Asn Ile Pro Thr Ile Phe Val Ser
          130            135            140

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ggc ggt ccg atg gaa gca ggt aaa gtt atc ggc gtg gca aac atc cag	480
Gly Gly Pro Met Glu Ala Gly Lys Val Ile Gly Val Ala Asn Ile Gln	
145 150 155 160	
ccc gaa cgc cgt ttg gac ttg att gac gcg atg att gaa tcg gcg gac	528
Pro Glu Arg Arg Leu Asp Leu Ile Asp Ala Met Ile Glu Ser Ala Asp	
165 170 175	
gac aat atc agc aac cgg caa gtt gaa gaa gtc gaa caa aac gcc tgc	576
Asp Asn Ile Ser Asn Arg Gln Val Glu Glu Val Glu Gln Asn Ala Cys	
180 185 190	
ccg acc tgc ggc tcg tgt tcg ggt atg ttt acg gca aac tcg atg aac	624
Pro Thr Cys Gly Ser Cys Ser Gly Met Phe Thr Ala Asn Ser Met Asn	
195 200 205	
tgc ctg acc gaa gca ctc ggg ctg tcc ctg ccg ggc aac ggt tct tac	672
Cys Leu Thr Glu Ala Leu Gly Leu Ser Leu Pro Gly Asn Gly Ser Tyr	
210 215 220	
ctc gcc acc cac gtc ggc cgc aaa gaa ctg ttc ctc gaa gcc ggc cgt	720
Leu Ala Thr His Val Gly Arg Lys Glu Leu Phe Leu Glu Ala Gly Arg	
225 230 235 240	
atg att gtc gaa atc acc aaa cgc tat tac gag caa aac gat gaa acc	768
Met Ile Val Glu Ile Thr Lys Arg Tyr Tyr Glu Gln Asn Asp Glu Thr	
245 250 255	
gtg tta ccg cgc agc att gcc acc aaa aaa gcg ttt gaa aac gct atg	816
Val Leu Pro Arg Ser Ile Ala Thr Lys Lys Ala Phe Glu Asn Ala Met	
260 265 270	
acc atg gac att gcc atg ggc ggc agc acc aac act att ttg cat tta	864
Thr Met Asp Ile Ala Met Gly Gly Ser Thr Asn Thr Ile Leu His Leu	
275 280 285	
ctc gcc gtt gcc aac gaa gcg ggc gtg gat ttc aaa atg gca gac atc	912
Leu Ala Val Ala Asn Glu Ala Gly Val Asp Phe Lys Met Ala Asp Ile	
290 295 300	
gac cgc tta agc cgc gtt gtg ccc tgc atc tgc aaa acc gca ccc aac	960
Asp Arg Leu Ser Arg Val Val Pro Cys Ile Cys Lys Thr Ala Pro Asn	
305 310 315 320	
aac cac gac tac tat atg gaa gac gtg cac cga gcc ggc ggc atc ttc	1008
Asn His Asp Tyr Tyr Met Glu Asp Val His Arg Ala Gly Gly Ile Phe	
325 330 335	

gcc atc ctg aaa gaa ttg gac aaa gcg ggc aag ctg cac acc gac gtg	1056
Ala Ile Leu Lys Glu Leu Asp Lys Ala Gly Lys Leu His Thr Asp Val	
340 345 350	
cac act atc cac gcg ccg acg ctg aaa gac gcg att gaa aaa tgg gac	1104
His Thr Ile His Ala Pro Thr Leu Lys Asp Ala Ile Glu Lys Trp Asp	
355 360 365	
gta acc aac cct gaa aac acc cac gcc atc gaa cgc ttc aaa gcc gcg	1152
Val Thr Asn Pro Glu Asn Thr His Ala Ile Glu Arg Phe Lys Ala Ala	
370 375 380	
ccc ggc ggc gta cgc acc acc caa gcg ttc tcg caa aac cgt atg tgg	1200
Pro Gly Gly Val Arg Thr Thr Gln Ala Phe Ser Gln Asn Arg Met Trp	
385 390 395 400	
aaa acc ctc gac ctc gac cgc gaa aaa ggc tgt atc cgc gac gtg gca	1248
Lys Thr Leu Asp Leu Asp Arg Glu Lys Gly Cys Ile Arg Asp Val Ala	
405 410 415	
cac gcc tac tcg caa gac ggc ggt ttg gcg gtc ttg ttc ggc aac atc	1296
His Ala Tyr Ser Gln Asp Gly Gly Leu Ala Val Leu Phe Gly Asn Ile	
420 425 430	
gcc gag cgc ggc tgc gtg gta aaa acc gca ggc gtg gac gag agc atc	1344
Ala Glu Arg Gly Cys Val Val Lys Thr Ala Gly Val Asp Glu Ser Ile	
435 440 445	
ctc aaa ttc acc ggc cgc gcc cgc gtg ttt gaa agc caa gaa gac gca	1392
Leu Lys Phe Thr Gly Arg Ala Arg Val Phe Glu Ser Gln Glu Asp Ala	
450 455 460	
gta gaa ggc att ttg ggc aac caa atc gtc gcc ggc gac atc gtc atc	1440
Val Glu Gly Ile Leu Gly Asn Gln Ile Val Ala Gly Asp Ile Val Ile	
465 470 475 480	
atc cgc tac gaa ggc ccg aaa ggc ggc ccg ggc atg cag gaa atg ctg	1488
Ile Arg Tyr Glu Gly Pro Lys Gly Gly Pro Gly Met Gln Glu Met Leu	
485 490 495	
tat ccg act tcc tac ctg aaa tct aaa ggc ctc ggc aaa gcc tgc gct	1536
Tyr Pro Thr Ser Tyr Leu Lys Ser Lys Gly Leu Gly Lys Ala Cys Ala	
500 505 510	
ctc tta acc gac gga cgc ttc tcc ggc ggc aca tca ggt ttg tcc atc	1584
Leu Leu Thr Asp Gly Arg Phe Ser Gly Gly Thr Ser Gly Leu Ser Ile	
515 520 525	

gga cac gcc tcg ccc gaa gcg gcg gaa ggt ggc gcg atc ggt ttg gta 1632
 Gly His Ala Ser Pro Glu Ala Ala Glu Gly Gly Ala Ile Gly Leu Val
 530 535 540

cac gaa ggc gat acc gtc gaa atc gac atc ccc aac cgc agc atc cgc 1680
 His Glu Gly Asp Thr Val Glu Ile Asp Ile Pro Asn Arg Ser Ile Arg
 545 550 555 560

ctt gtc att tcc gac gaa gaa ctc gcc gcc cgc cgt gcc gaa atg gaa 1728
 Leu Val Ile Ser Asp Glu Glu Leu Ala Ala Arg Arg Ala Glu Met Glu
 565 570 575

gca cgc ggc agc aaa gcg tgg aaa ccg gaa aac cgc gac cgc tac gtc 1776
 Ala Arg Gly Ser Lys Ala Trp Lys Pro Glu Asn Arg Asp Arg Tyr Val
 580 585 590

tcc gcc gca tta aga gct tac ggc gcg atg gcg act tcc gcc gac aaa 1824
 Ser Ala Ala Leu Arg Ala Tyr Gly Ala Met Ala Thr Ser Ala Asp Lys
 595 600 605

ggc gcg gtg cgc gac gta gcg caa atc gaa aga 1857
 Gly Ala Val Arg Asp Val Ala Gln Ile Glu Arg
 610 615

<210> 28

<211> 619

<212> PRT

<213> Neisseria meningitidis

<400> 28

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Gly Ala Arg Ala Leu Trp Arg Ala Thr Gly Val Met Glu Thr Asp Phe
 20 25 30

Gly Lys Pro Ile Ile Ala Val Ala Asn Ser Phe Thr Gln Phe Val Pro
 35 40 45

Gly His Val His Leu His Asn Met Gly Gln Leu Val Ala Arg Glu Ile
 50 55 60

Glu Lys Ala Gly Ala Ile Ala Lys Glu Phe Asn Thr Ile Ala Ile Asp
 65 70 75 80

Asp Gly Ile Ala Met Gly His Ser Gly Met Leu Tyr Ser Leu Pro Ser
 85 90 95

Arg Asp Leu Ile Ala Asp Ser Ile Glu Tyr Met Val Asn Ala His Cys
 100 105 110

Ala Asp Ala Leu Val Cys Ile Ser Asn Cys Asp Lys Ile Thr Pro Gly
 115 120 125

Met Leu Ile Ala Ala Met Arg Leu Asn Ile Pro Thr Ile Phe Val Ser
 130 135 140

Gly Gly Pro Met Glu Ala Gly Lys Val Ile Gly Val Ala Asn Ile Gln
 145 150 155 160

Pro Glu Arg Arg Leu Asp Leu Ile Asp Ala Met Ile Glu Ser Ala Asp
 165 170 175

Asp Asn Ile Ser Asn Arg Gln Val Glu Glu Val Glu Gln Asn Ala Cys
 180 185 190

Pro Thr Cys Gly Ser Cys Ser Gly Met Phe Thr Ala Asn Ser Met Asn
 195 200 205

Cys Leu Thr Glu Ala Leu Gly Leu Ser Leu Pro Gly Asn Gly Ser Tyr
 210 215 220

Leu Ala Thr His Val Gly Arg Lys Glu Leu Phe Leu Glu Ala Gly Arg
 225 230 235 240

Met Ile Val Glu Ile Thr Lys Arg Tyr Tyr Glu Gln Asn Asp Glu Thr
 245 250 255

Val Leu Pro Arg Ser Ile Ala Thr Lys Lys Ala Phe Glu Asn Ala Met
 260 265 270

Thr Met Asp Ile Ala Met Gly Gly Ser Thr Asn Thr Ile Leu His Leu
 275 280 285

Leu Ala Val Ala Asn Glu Ala Gly Val Asp Phe Lys Met Ala Asp Ile
 290 295 300

Asp Arg Leu Ser Arg Val Val Pro Cys Ile Cys Lys Thr Ala Pro Asn
 305 310 315 320

Asn His Asp Tyr Tyr Met Glu Asp Val His Arg Ala Gly Gly Ile Phe
 325 330 335

Ala Ile Leu Lys Glu Leu Asp Lys Ala Gly Lys Leu His Thr Asp Val
 340 345 350

His Thr Ile His Ala Pro Thr Leu Lys Asp Ala Ile Glu Lys Trp Asp
 355 360 365
 Val Thr Asn Pro Glu Asn Thr His Ala Ile Glu Arg Phe Lys Ala Ala
 370 375 380
 Pro Gly Gly Val Arg Thr Thr Gln Ala Phe Ser Gln Asn Arg Met Trp
 385 390 395 400
 Lys Thr Leu Asp Leu Asp Arg Glu Lys Gly Cys Ile Arg Asp Val Ala
 405 410 415
 His Ala Tyr Ser Gln Asp Gly Gly Leu Ala Val Leu Phe Gly Asn Ile
 420 425 430
 Ala Glu Arg Gly Cys Val Val Lys Thr Ala Gly Val Asp Glu Ser Ile
 435 440 445
 Leu Lys Phe Thr Gly Arg Ala Arg Val Phe Glu Ser Gln Glu Asp Ala
 450 455 460
 Val Glu Gly Ile Leu Gly Asn Gln Ile Val Ala Gly Asp Ile Val Ile
 465 470 475 480
 Ile Arg Tyr Glu Gly Pro Lys Gly Gly Pro Gly Met Gln Glu Met Leu
 485 490 495
 Tyr Pro Thr Ser Tyr Leu Lys Ser Lys Gly Leu Gly Lys Ala Cys Ala
 500 505 510
 Leu Leu Thr Asp Gly Arg Phe Ser Gly Gly Thr Ser Gly Leu Ser Ile
 515 520 525
 Gly His Ala Ser Pro Glu Ala Ala Glu Gly Gly Ala Ile Gly Leu Val
 530 535 540
 His Glu Gly Asp Thr Val Glu Ile Asp Ile Pro Asn Arg Ser Ile Arg
 545 550 555 560
 Leu Val Ile Ser Asp Glu Glu Leu Ala Ala Arg Arg Ala Glu Met Glu
 565 570 575
 Ala Arg Gly Ser Lys Ala Trp Lys Pro Glu Asn Arg Asp Arg Tyr Val
 580 585 590
 Ser Ala Ala Leu Arg Ala Tyr Gly Ala Met Ala Thr Ser Ala Asp Lys
 595 600 605

Gly Ala Val Arg Asp Val Ala Gln Ile Glu Arg
 610 615

<210> 29

<211> 1725

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(1725)

<400> 29

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 1 5 10 15

ggc gtc gag tac gtt ttc ggt tat ccc ggc ggt gcg gtt atc gaa atc 96
 Gly Val Glu Tyr Val Phe Gly Tyr Pro Gly Gly Ala Val Ile Glu Ile
 20 25 30

tac gat gcc ctt ttc caa ctc aat aaa ttc aag cac att ctg acc cgt 144
 Tyr Asp Ala Leu Phe Gln Leu Asn Lys Phe Lys His Ile Leu Thr Arg
 35 40 45

cac gag cag gcg gca gta cac gcg gca gat gcg tat gcg cgc gtc agc 192
 His Glu Gln Ala Ala Val His Ala Ala Asp Ala Tyr Ala Arg Val Ser
 50 55 60

ggc aag gtg ggc gtg gca ttg gtt aca tcc ggc ccg ggc gtt acc aat 240
 Gly Lys Val Gly Val Ala Leu Val Thr Ser Gly Pro Gly Val Thr Asn
 65 70 75 80

gca ctg acc ggt att gct act gcc tat acg gat tcg att ccg atg gtg 288
 Ala Leu Thr Gly Ile Ala Thr Ala Tyr Thr Asp Ser Ile Pro Met Val
 85 90 95

gtc atc agc ggg cag gta ggc aat tcc ctg att ggt acg gat gcg ttc 336
 Val Ile Ser Gly Gln Val Gly Asn Ser Leu Ile Gly Thr Asp Ala Phe
 100 105 110

caa gaa gtt gat acg gtg ggt att acc cgt ccg tgc gtc aaa cac aat 384
 Gln Glu Val Asp Thr Val Gly Ile Thr Arg Pro Cys Val Lys His Asn
 115 120 125


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ttc ctg gtt acg gac atc aat gag ttg gcg gaa acc att aaa aag gcg 432
Phe Leu Val Thr Asp Ile Asn Glu Leu Ala Glu Thr Ile Lys Lys Ala
130 135 140

ttc caa att gcc gca agc ggc cga ccg ggg ccc gtg gtg gtt gat gtc 480
Phe Gln Ile Ala Ala Ser Gly Arg Pro Gly Pro Val Val Val Asp Val
145 150 155 160

ccg aaa gat gtt acg cag gcg atg gcg aaa ttc agc tat cct cag gaa 528
Pro Lys Asp Val Thr Gln Ala Met Ala Lys Phe Ser Tyr Pro Gln Glu
165 170 175

gac att ttt atc cgt tcg tat caa ccg gtt gtt caa ggg cat atc ggt 576
Asp Ile Phe Ile Arg Ser Tyr Gln Pro Val Val Gln Gly His Ile Gly
180 185 190

cag att aaa aag gcc gtg cag atg ttg gca tct gcc aaa cgt ccg gtc 624
Gln Ile Lys Lys Ala Val Gln Met Leu Ala Ser Ala Lys Arg Pro Val
195 200 205

gtc tat ttt ggc ggc ggc gtg gtg ttg ggt aat gct tct gaa gag ctg 672
Val Tyr Phe Gly Gly Gly Val Val Leu Gly Asn Ala Ser Glu Glu Leu
210 215 220

acc cga ttt gtc cga atg acg ggt gct ccg tgc acg ggt acg ctg atg 720
Thr Arg Phe Val Arg Met Thr Gly Ala Pro Cys Thr Gly Thr Leu Met
225 230 235 240

ggc ttg ggc gct tat cct tcc ggc gac cgc caa ttc ctc ggt atg ctc 768
Gly Leu Gly Ala Tyr Pro Ser Gly Asp Arg Gln Phe Leu Gly Met Leu
245 250 255

ggt atg cac ggt act tac gag gca aac ctt gcc atg cag aat gcg gat 816
Gly Met His Gly Thr Tyr Glu Ala Asn Leu Ala Met Gln Asn Ala Asp
260 265 270

gtc gtt ctt gcc gta ggt gcg cgt ttt gac gac cgt gtg gtt tcc gta 864
Val Val Leu Ala Val Gly Ala Arg Phe Asp Asp Arg Val Val Ser Val
275 280 285

ccg tcc aaa ttt ttc gag aag gcg aag aag gtt atc cat atc gat gtc 912
Pro Ser Lys Phe Phe Glu Lys Ala Lys Lys Val Ile His Ile Asp Val
290 295 300

gat cct tcc agc atc gcc aaa cgc gtg aag gcg gac att ccg att gtc 960
Asp Pro Ser Ser Ile Ala Lys Arg Val Lys Ala Asp Ile Pro Ile Val
305 310 315 320

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ggc gac gtg aaa aac gtt ttg tcc gag atg gtt gcg ctg tgg caa aaa 1008
Gly Asp Val Lys Asn Val Leu Ser Glu Met Val Ala Leu Trp Gln Lys
          325                330                335

caa gag tcc gtg ccg tct gaa gat gct ttg ggc aaa tgg tgg aaa acc 1056
Gln Glu Ser Val Pro Ser Glu Asp Ala Leu Gly Lys Trp Trp Lys Thr
          340                345                350

ata gag gaa tgg cgt tcc cgc gat tgc ttg tgg ttt gac aac ggc agc 1104
Ile Glu Glu Trp Arg Ser Arg Asp Cys Leu Trp Phe Asp Asn Gly Ser
          355                360                365

gaa att atc aag cca caa tat gtg att cag aag ctt gcc gag att acc 1152
Glu Ile Ile Lys Pro Gln Tyr Val Ile Gln Lys Leu Ala Glu Ile Thr
          370                375                380

ggc aat tcg gca atc atc aca tcg gat gta ggg cag cat caa atg ttt 1200
Gly Asn Ser Ala Ile Ile Thr Ser Asp Val Gly Gln His Gln Met Phe
          385                390                395                400

gcg gct caa tat tat ccc ttc gaa cgt ccg cgc caa tgg ctc aac tcc 1248
Ala Ala Gln Tyr Tyr Pro Phe Glu Arg Pro Arg Gln Trp Leu Asn Ser
          405                410                415

ggc ggt ttg ggt acg atg ggc gtt ggt ctg cct tat gcg att ggt gca 1296
Gly Gly Leu Gly Thr Met Gly Val Gly Leu Pro Tyr Ala Ile Gly Ala
          420                425                430

aaa ctt gcc gcc ccg gat caa gac gta ttc tgt att acc ggc gac ggc 1344
Lys Leu Ala Ala Pro Asp Gln Asp Val Phe Cys Ile Thr Gly Asp Gly
          435                440                445

tcg atc cag atg aac atc caa gag ctg tcc acc tgt ttc caa tat cgg 1392
Ser Ile Gln Met Asn Ile Gln Glu Leu Ser Thr Cys Phe Gln Tyr Arg
          450                455                460

att ccg gtt aac gtc att acg ctg aac aac ggt tat ctc ggt atg gta 1440
Ile Pro Val Asn Val Ile Thr Leu Asn Asn Gly Tyr Leu Gly Met Val
          465                470                475                480

cgc cag tgg cag gaa ata tat tac ggc gga cga gag tcg gaa acc tat 1488
Arg Gln Trp Gln Glu Ile Tyr Tyr Gly Gly Arg Glu Ser Glu Thr Tyr
          485                490                495

ttc gat tct ttg ccc gat ttc gtc aaa ctt gcc gag gca tac ggc cat 1536
Phe Asp Ser Leu Pro Asp Phe Val Lys Leu Ala Glu Ala Tyr Gly His
          500                505                510

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atc ggt atc cgc gtg gac aag aag tct gat gtg gaa ggt gcg ttg ttg 1584
 Ile Gly Ile Arg Val Asp Lys Lys Ser Asp Val Glu Gly Ala Leu Leu
 515 520 525

gaa gca ttg aac caa aaa gac agg ctg gtg ttt atc gac ttc ctg acc 1632
 Glu Ala Leu Asn Gln Lys Asp Arg Leu Val Phe Ile Asp Phe Leu Thr
 530 535 540

gac cag aaa cag aat gtg atg ccc atg gtc ggc aac ggc aaa ggt ttg 1680
 Asp Gln Lys Gln Asn Val Met Pro Met Val Gly Asn Gly Lys Gly Leu
 545 550 555 560

gac gaa atg gta ctt ccg ccg cat atg cgt gcg gac gga aag gcg 1725
 Asp Glu Met Val Leu Pro Pro His Met Arg Ala Asp Gly Lys Ala
 565 570 575

<210> 30

<211> 575

<212> PRT

<213> Neisseria meningitidis

<400> 30

Met Gln Leu Ser Gly Ala Gln Ile Ile Val Gln Ser Leu Lys Ala Glu
 1 5 10 15

Gly Val Glu Tyr Val Phe Gly Tyr Pro Gly Gly Ala Val Ile Glu Ile
 20 25 30

Tyr Asp Ala Leu Phe Gln Leu Asn Lys Phe Lys His Ile Leu Thr Arg
 35 40 45

His Glu Gln Ala Ala Val His Ala Ala Asp Ala Tyr Ala Arg Val Ser
 50 55 60

Gly Lys Val Gly Val Ala Leu Val Thr Ser Gly Pro Gly Val Thr Asn
 65 70 75 80

Ala Leu Thr Gly Ile Ala Thr Ala Tyr Thr Asp Ser Ile Pro Met Val
 85 90 95

Val Ile Ser Gly Gln Val Gly Asn Ser Leu Ile Gly Thr Asp Ala Phe
 100 105 110

Gln Glu Val Asp Thr Val Gly Ile Thr Arg Pro Cys Val Lys His Asn
 115 120 125

Phe Leu Val Thr Asp Ile Asn Glu Leu Ala Glu Thr Ile Lys Lys Ala

130	135	140
Phe Gln Ile Ala Ala Ser Gly Arg Pro Gly Pro Val Val Val Asp Val		
145	150	155 160
Pro Lys Asp Val Thr Gln Ala Met Ala Lys Phe Ser Tyr Pro Gln Glu		
	165	170 175
Asp Ile Phe Ile Arg Ser Tyr Gln Pro Val Val Gln Gly His Ile Gly		
	180	185 190
Gln Ile Lys Lys Ala Val Gln Met Leu Ala Ser Ala Lys Arg Pro Val		
	195	200 205
Val Tyr Phe Gly Gly Gly Val Val Leu Gly Asn Ala Ser Glu Glu Leu		
	210	215 220
Thr Arg Phe Val Arg Met Thr Gly Ala Pro Cys Thr Gly Thr Leu Met		
	225	230 235 240
Gly Leu Gly Ala Tyr Pro Ser Gly Asp Arg Gln Phe Leu Gly Met Leu		
	245	250 255
Gly Met His Gly Thr Tyr Glu Ala Asn Leu Ala Met Gln Asn Ala Asp		
	260	265 270
Val Val Leu Ala Val Gly Ala Arg Phe Asp Asp Arg Val Val Ser Val		
	275	280 285
Pro Ser Lys Phe Phe Glu Lys Ala Lys Lys Val Ile His Ile Asp Val		
	290	295 300
Asp Pro Ser Ser Ile Ala Lys Arg Val Lys Ala Asp Ile Pro Ile Val		
	305	310 315 320
Gly Asp Val Lys Asn Val Leu Ser Glu Met Val Ala Leu Trp Gln Lys		
	325	330 335
Gln Glu Ser Val Pro Ser Glu Asp Ala Leu Gly Lys Trp Trp Lys Thr		
	340	345 350
Ile Glu Glu Trp Arg Ser Arg Asp Cys Leu Trp Phe Asp Asn Gly Ser		
	355	360 365
Glu Ile Ile Lys Pro Gln Tyr Val Ile Gln Lys Leu Ala Glu Ile Thr		
	370	375 380
Gly Asn Ser Ala Ile Ile Thr Ser Asp Val Gly Gln His Gln Met Phe		

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385              390              395              400

Ala Ala Gln Tyr Tyr Pro Phe Glu Arg Pro Arg Gln Trp Leu Asn Ser
              405              410              415

Gly Gly Leu Gly Thr Met Gly Val Gly Leu Pro Tyr Ala Ile Gly Ala
              420              425              430

Lys Leu Ala Ala Pro Asp Gln Asp Val Phe Cys Ile Thr Gly Asp Gly
              435              440              445

Ser Ile Gln Met Asn Ile Gln Glu Leu Ser Thr Cys Phe Gln Tyr Arg
              450              455              460

Ile Pro Val Asn Val Ile Thr Leu Asn Asn Gly Tyr Leu Gly Met Val
465              470              475              480

Arg Gln Trp Gln Glu Ile Tyr Tyr Gly Gly Arg Glu Ser Glu Thr Tyr
              485              490              495

Phe Asp Ser Leu Pro Asp Phe Val Lys Leu Ala Glu Ala Tyr Gly His
              500              505              510

Ile Gly Ile Arg Val Asp Lys Lys Ser Asp Val Glu Gly Ala Leu Leu
              515              520              525

Glu Ala Leu Asn Gln Lys Asp Arg Leu Val Phe Ile Asp Phe Leu Thr
              530              535              540

Asp Gln Lys Gln Asn Val Met Pro Met Val Gly Asn Gly Lys Gly Leu
545              550              555              560

Asp Glu Met Val Leu Pro Pro His Met Arg Ala Asp Gly Lys Ala
              565              570              575

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<210> 31

<211> 651

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(651)

<400> 31

atg ttt tcc gta ccg cgt tcc ttt ttg ccg ggc gtt ttc gta ctt gcc 48

Met	Phe	Ser	Val	Pro	Arg	Ser	Phe	Leu	Pro	Gly	Val	Phe	Val	Leu	Ala	
1				5					10					15		
gcg	ctt	gcc	gcc	tgc	aaa	cct	caa	gac	aac	agt	gcg	gcg	caa	gtc	gct	96
Ala	Leu	Ala	Ala	Cys	Lys	Pro	Gln	Asp	Asn	Ser	Ala	Ala	Gln	Val	Ala	
		20						25					30			
tct	tca	agt	gca	tcc	gcg	tcg	gct	gcg	gaa	aat	gcg	gca	aag	ccg	caa	144
Ser	Ser	Ser	Ala	Ser	Ala	Ser	Ala	Ala	Glu	Asn	Ala	Ala	Lys	Pro	Gln	
		35					40					45				
acg	cgc	ggt	acg	gat	atg	cgt	aag	gaa	gac	atc	ggc	ggc	gat	ttc	acg	192
Thr	Arg	Gly	Thr	Asp	Met	Arg	Lys	Glu	Asp	Ile	Gly	Gly	Asp	Phe	Thr	
		50				55				60						
ctg	acc	gac	ggc	gaa	ggc	aag	cct	ttc	aac	ctg	agc	gat	ttg	aaa	ggc	240
Leu	Thr	Asp	Gly	Glu	Gly	Lys	Pro	Phe	Asn	Leu	Ser	Asp	Leu	Lys	Gly	
65					70					75				80		
aag	gtc	gtg	att	ctg	tct	ttc	ggc	ttt	acg	cac	tgt	ccc	gat	gtc	tgc	288
Lys	Val	Val	Ile	Leu	Ser	Phe	Gly	Phe	Thr	His	Cys	Pro	Asp	Val	Cys	
			85					90					95			
ccg	aca	gag	ctt	ttg	acg	tac	agc	gac	acg	ttg	aag	cag	ttg	ggc	ggg	336
Pro	Thr	Glu	Leu	Leu	Thr	Tyr	Ser	Asp	Thr	Leu	Lys	Gln	Leu	Gly	Gly	
		100					105					110				
cag	gct	aag	gac	gtg	aaa	gtg	gtg	ttc	gtc	agc	atc	gat	ccg	gaa	cgc	384
Gln	Ala	Lys	Asp	Val	Lys	Val	Val	Phe	Val	Ser	Ile	Asp	Pro	Glu	Arg	
		115				120					125					
gac	acg	cct	gaa	atc	atc	ggc	aag	tat	gcc	aaa	cag	ttc	aat	ccg	gac	432
Asp	Thr	Pro	Glu	Ile	Ile	Gly	Lys	Tyr	Ala	Lys	Gln	Phe	Asn	Pro	Asp	
		130				135				140						
ttt	atc	ggt	ctg	acg	gca	acg	ggc	gac	caa	aac	ctg	ccg	gtc	atc	aag	480
Phe	Ile	Gly	Leu	Thr	Ala	Thr	Gly	Asp	Gln	Asn	Leu	Pro	Val	Ile	Lys	
145				150				155				160				
cag	caa	tac	cgc	gtg	gtt	tct	gcc	aaa	gtc	aat	caa	aaa	gac	gac	agc	528
Gln	Gln	Tyr	Arg	Val	Val	Ser	Ala	Lys	Val	Asn	Gln	Lys	Asp	Asp	Ser	
			165				170				175					
gaa	aac	tat	ttg	gtc	gac	cac	tct	tcc	ggt	gcg	tat	ctc	atc	gac	aaa	576
Glu	Asn	Tyr	Leu	Val	Asp	His	Ser	Ser	Gly	Ala	Tyr	Leu	Ile	Asp	Lys	
		180					185				190					
aac	ggt	gag	gtt	gcc	att	ttc	tcg	cct	tac	gga	agc	gag	ccg	gaa	acg	624

Asn Gly Glu Val Ala Ile Phe Ser Pro Tyr Gly Ser Glu Pro Glu Thr
 195 200 205

att gct gcc gat gta agg acc ctg ctc 651
 ile Ala Ala Asp Val Arg Thr Leu Leu
 210 215

<210> 32

<211> 217

<212> PRT

<213> Neisseria meningitidis

<400> 32

Met Phe Ser Val Pro Arg Ser Phe Leu Pro Gly Val Phe Val Leu Ala
 1 5 10 15

Ala Leu Ala Ala Cys Lys Pro Gln Asp Asn Ser Ala Ala Gln Val Ala
 20 25 30

Ser Ser Ser Ala Ser Ala Ser Ala Ala Glu Asn Ala Ala Lys Pro Gln
 35 40 45

Thr Arg Gly Thr Asp Met Arg Lys Glu Asp Ile Gly Gly Asp Phe Thr
 50 55 60

Leu Thr Asp Gly Glu Gly Lys Pro Phe Asn Leu Ser Asp Leu Lys Gly
 65 70 75 80

Lys Val Val Ile Leu Ser Phe Gly Phe Thr His Cys Pro Asp Val Cys
 85 90 95

Pro Thr Glu Leu Leu Thr Tyr Ser Asp Thr Leu Lys Gln Leu Gly Gly
 100 105 110

Gln Ala Lys Asp Val Lys Val Val Phe Val Ser Ile Asp Pro Glu Arg
 115 120 125

Asp Thr Pro Glu Ile Ile Gly Lys Tyr Ala Lys Gln Phe Asn Pro Asp
 130 135 140

Phe Ile Gly Leu Thr Ala Thr Gly Asp Gln Asn Leu Pro Val Ile Lys
 145 150 155 160

Gln Gln Tyr Arg Val Val Ser Ala Lys Val Asn Gln Lys Asp Asp Ser
 165 170 175

Glu Asn Tyr Leu Val Asp His Ser Ser Gly Ala Tyr Leu Ile Asp Lys

[illegible]

<210> 34

<211> 110

<212> PRT

<213> Neisseria meningitidis

<400> 34

Leu Pro Leu Leu Cys Cys Leu Trp Phe Trp Gln Pro Ala Ala Val Lys
 1 5 10 15

Lys Pro Leu Lys Leu Pro Leu Leu Lys His Leu Pro Pro Lys Leu Pro
 20 25 30

Leu Leu Lys His Leu Pro Pro Lys Leu Pro Leu Leu Lys His Leu Pro
 35 40 45

Pro Lys Leu Leu Leu Leu Lys His Leu Pro Pro Lys Leu Pro Leu Leu
 50 55 60

Lys Leu Pro Leu Pro Lys His Leu Pro Leu Lys Leu Pro Leu Pro Lys
 65 70 75 80

His Leu Pro Leu Lys Leu Leu Leu Pro Lys Leu Gln Asn Lys His Phe
 85 90 95

Pro Leu Ala Lys Lys Gln Asp Thr Phe Ser Ile Leu Leu Phe
 100 105 110

<210> 35

<211> 2118

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(2118)

<400> 35

atg ttc gac aaa cac gtt aag acc ttc caa tac ggt aat cag acc gtt 48
 Met Phe Asp Lys His Val Lys Thr Phe Gln Tyr Gly Asn Gln Thr Val
 1 5 10 15

act ttg gaa acc ggc gaa att gcc cgc caa gcc gcc gct gcc gtt aaa 96
 Thr Leu Glu Thr Gly Glu Ile Ala Arg Gln Ala Ala Ala Ala Val Lys
 20 25 30

gtc tct atg ggc gac acc gtg gtt ttg gtt gcc gtt acc acc aac aaa	144
Val Ser Met Gly Asp Thr Val Val Leu Val Ala Val Thr Thr Asn Lys	
35 40 45	
gaa gtg aaa gaa ggc caa gac ttc ttc ccc ctg acc gtc gat tat ttg	192
Glu Val Lys Glu Gly Gln Asp Phe Phe Pro Leu Thr Val Asp Tyr Leu	
50 55 60	
gaa cgc act tac gcc gca ggt aaa atc ccc ggc ggt ttc ttc aaa cgc	240
Glu Arg Thr Tyr Ala Ala Gly Lys Ile Pro Gly Gly Phe Phe Lys Arg	
65 70 75 80	
gaa ggc aaa caa agc gaa aaa gaa atc ctg acc agc cgt ctg atc gac	288
Glu Gly Lys Gln Ser Glu Lys Glu Ile Leu Thr Ser Arg Leu Ile Asp	
85 90 95	
cgc ccg att cgt ccg ctg ttc ccc gaa ggt ttc tac cac gac atc caa	336
Arg Pro Ile Arg Pro Leu Phe Pro Glu Gly Phe Tyr His Asp Ile Gln	
100 105 110	
atc gta gcg atg gtc gtg tcc gtc gat ccc gaa atc gat tcc gac att	384
Ile Val Ala Met Val Val Ser Val Asp Pro Glu Ile Asp Ser Asp Ile	
115 120 125	
cct gca atg ttg ggc gcg tct gcc gcg ctg gtg ttg agc ggc gta ccg	432
Pro Ala Met Leu Gly Ala Ser Ala Ala Leu Val Leu Ser Gly Val Pro	
130 135 140	
ttt gcc ggt ccg att ggt gcg gca cgc gtg ggt tat ata aac ggc gtg	480
Phe Ala Gly Pro Ile Gly Ala Ala Arg Val Gly Tyr Ile Asn Gly Val	
145 150 155 160	
tac gtt ttg aac ccg act aaa gcc gaa ttg gcg aaa tcg caa ttg gac	528
Tyr Val Leu Asn Pro Thr Lys Ala Glu Leu Ala Lys Ser Gln Leu Asp	
165 170 175	
ttg gtg gtc gcc ggt act tca aaa gcc gtg ttg atg gtg gaa tcc gaa	576
Leu Val Val Ala Gly Thr Ser Lys Ala Val Leu Met Val Glu Ser Glu	
180 185 190	
gcc aaa atc ctg ccc gaa gac gtg atg ctg ggc gcg gtg gtt tac ggc	624
Ala Lys Ile Leu Pro Glu Asp Val Met Leu Gly Ala Val Val Tyr Gly	
195 200 205	
cac gat caa atg cag gtt gcc atc aat gca atc aat gaa ttt gcc gac	672
His Asp Gln Met Gln Val Ala Ile Asn Ala Ile Asn Glu Phe Ala Asp	
210 215 220	

65

ccg aaa cct gaa gat ttc agc tac acc atg cgc gtg gtc tcc gaa att	1296
Pro Lys Pro Glu Asp Phe Ser Tyr Thr Met Arg Val Val Ser Glu Ile	
420 425 430	
acc gaa tcc aac ggc tct tcc tct atg gct tcc gtc tgc ggc ggc tgc	1344
Thr Glu Ser Asn Gly Ser Ser Ser Met Ala Ser Val Cys Gly Gly Cys	
435 440 445	
ttg agc ctg ctg tcc gcc ggc gtg cct ttg aaa gca cac gtt gcc ggt	1392
Leu Ser Leu Leu Ser Ala Gly Val Pro Leu Lys Ala His Val Ala Gly	
450 455 460	
atc gcg atg ggt ctg att ctg gaa ggc aat aag ttt gcc gtc ctg acc	1440
Ile Ala Met Gly Leu Ile Leu Glu Gly Asn Lys Phe Ala Val Leu Thr	
465 470 475 480	
gac att ttg ggc gac gaa gac cac ttg ggc gat atg gac ttt aaa gtg	1488
Asp Ile Leu Gly Asp Glu Asp His Leu Gly Asp Met Asp Phe Lys Val	
485 490 495	
gct ggt acg acc gaa ggc gtt acc gcg ctg caa atg gac atc aaa atc	1536
Ala Gly Thr Thr Glu Gly Val Thr Ala Leu Gln Met Asp Ile Lys Ile	
500 505 510	
caa ggc att acc aaa gag att atg caa atc gct ttg gcg cag gcc aaa	1584
Gln Gly Ile Thr Lys Glu Ile Met Gln Ile Ala Leu Ala Gln Ala Lys	
515 520 525	
gaa gcg cgt ctg cac atc ttg gat cag atg aaa gcc gcc gtt gcg ggc	1632
Glu Ala Arg Leu His Ile Leu Asp Gln Met Lys Ala Ala Val Ala Gly	
530 535 540	
ccg caa gag ctg tcc gca cac gcg cca cgc ttg ttc acg atg aaa atc	1680
Pro Gln Glu Leu Ser Ala His Ala Pro Arg Leu Phe Thr Met Lys Ile	
545 550 555 560	
aac caa gac aaa atc cgc gaa gtt atc ggt aag ggc ggt gaa acc atc	1728
Asn Gln Asp Lys Ile Arg Glu Val Ile Gly Lys Gly Gly Glu Thr Ile	
565 570 575	
cgt gcg att acc gct gaa acc ggt acg gaa atc aat att gcc gaa gac	1776
Arg Ala Ile Thr Ala Glu Thr Gly Thr Glu Ile Asn Ile Ala Glu Asp	
580 585 590	
ggt acg att acc att gcc gca acc act caa gaa gcc ggc gat gcg gcg	1824
Gly Thr Ile Thr Ile Ala Ala Thr Thr Gln Glu Ala Gly Asp Ala Ala	
595 600 605	

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aaa aaa cgc atc gag cag att act gcc gaa gtg gaa gtg ggc aaa gtg 1872
Lys Lys Arg Ile Glu Gln Ile Thr Ala Glu Val Glu Val Gly Lys Val
    610                615                620

tac gaa ggc act gtg gtg aaa atc ctc gac aac aat gtc ggc gcg att 1920
Tyr Glu Gly Thr Val Val Lys Ile Leu Asp Asn Asn Val Gly Ala Ile
625                630                635                640

gtc agt gtg atg ccg ggc aaa gac ggt ttg gta cac atc agc caa atc 1968
Val Ser Val Met Pro Gly Lys Asp Gly Leu Val His Ile Ser Gln Ile
                645                650                655

gcc cac gag cgc gta cgc aat gtc ggc gac tac ctg caa gtc ggt cag 2016
Ala His Glu Arg Val Arg Asn Val Gly Asp Tyr Leu Gln Val Gly Gln
                660                665                670

gtg gtg aac gtg aaa gca ttg gaa gtg gac gac aga ggc cgt gtc cgt 2064
Val Val Asn Val Lys Ala Leu Glu Val Asp Asp Arg Gly Arg Val Arg
                675                680                685

ctg tcc atc aaa gcc ctg ctg gac gcg cct gcc cgt gag gaa aat gcc 2112
Leu Ser Ile Lys Ala Leu Leu Asp Ala Pro Ala Arg Glu Glu Asn Ala
    690                695                700

gcc gag
Ala Glu
705

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<210> 36

<211> 706

<212> PRT

<213> Neisseria meningitidis

<400> 36

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Met Phe Asp Lys His Val Lys Thr Phe Gln Tyr Gly Asn Gln Thr Val
    1                5                10                15

Thr Leu Glu Thr Gly Glu Ile Ala Arg Gln Ala Ala Ala Val Lys
    20                25                30

Val Ser Met Gly Asp Thr Val Val Leu Val Ala Val Thr Thr Asn Lys
    35                40                45

Glu Val Lys Glu Gly Gln Asp Phe Phe Pro Leu Thr Val Asp Tyr Leu
    50                55                60

Glu Arg Thr Tyr Ala Ala Gly Lys Ile Pro Gly Gly Phe Phe Lys Arg

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65		70		75		80
Glu Gly Lys Gln Ser Glu Lys Glu Ile Leu Thr Ser Arg Leu Ile Asp						
		85		90		95
Arg Pro Ile Arg Pro Leu Phe Pro Glu Gly Phe Tyr His Asp Ile Gln						
	100		105		110	
Ile Val Ala Met Val Val Ser Val Asp Pro Glu Ile Asp Ser Asp Ile						
	115		120		125	
Pro Ala Met Leu Gly Ala Ser Ala Ala Leu Val Leu Ser Gly Val Pro						
	130		135		140	
Phe Ala Gly Pro Ile Gly Ala Ala Arg Val Gly Tyr Ile Asn Gly Val						
145		150		155		160
Tyr Val Leu Asn Pro Thr Lys Ala Glu Leu Ala Lys Ser Gln Leu Asp						
	165		170		175	
Leu Val Val Ala Gly Thr Ser Lys Ala Val Leu Met Val Glu Ser Glu						
	180		185		190	
Ala Lys Ile Leu Pro Glu Asp Val Met Leu Gly Ala Val Val Tyr Gly						
	195		200		205	
His Asp Gln Met Gln Val Ala Ile Asn Ala Ile Asn Glu Phe Ala Asp						
	210		215		220	
Glu Val Asn Pro Glu Leu Trp Asp Trp Lys Ala Pro Glu Thr Asn Glu						
225		230		235		240
Glu Leu Val Ala Lys Val Arg Gly Ile Ala Gly Glu Thr Ile Lys Glu						
	245		250		255	
Ala Phe Lys Ile Arg Gln Lys Gln Ala Arg Ser Ala Lys Leu Asp Glu						
	260		265		270	
Ala Trp Ser Ala Val Lys Glu Ala Leu Ile Thr Glu Glu Thr Asp Thr						
	275		280		285	
Leu Ala Ala Asn Glu Ile Lys Gly Ile Phe Lys His Leu Glu Ala Asp						
	290		295		300	
Val Val Arg Ser Gln Ile Leu Asp Gly Gln Pro Arg Ile Asp Gly Arg						
305		310		315		320
Asp Thr Arg Thr Val Arg Pro Leu Asn Ile Gln Thr Gly Val Leu Pro						

	325		330		335
Arg Thr His Gly Ser Ala Leu Phe Thr Arg Gly Glu Thr Gln Ala Leu					
	340		345		350
Ala Val Ala Thr Leu Gly Thr Ser Arg Asp Glu Gln Ile Ile Asp Ala					
	355		360		365
Leu Ser Gly Glu Tyr Thr Asp Arg Phe Met Leu His Tyr Asn Phe Pro					
	370		375		380
Pro Tyr Ser Thr Gly Glu Val Gly Arg Met Gly Ala Pro Lys Arg Arg					
385		390		395	400
Glu Ile Gly His Gly Arg Leu Ala Lys Arg Ala Leu Leu Ala Val Leu					
	405		410		415
Pro Lys Pro Glu Asp Phe Ser Tyr Thr Met Arg Val Val Ser Glu Ile					
	420		425		430
Thr Glu Ser Asn Gly Ser Ser Ser Met Ala Ser Val Cys Gly Gly Cys					
	435		440		445
Leu Ser Leu Leu Ser Ala Gly Val Pro Leu Lys Ala His Val Ala Gly					
	450		455		460
Ile Ala Met Gly Leu Ile Leu Glu Gly Asn Lys Phe Ala Val Leu Thr					
465		470		475	480
Asp Ile Leu Gly Asp Glu Asp His Leu Gly Asp Met Asp Phe Lys Val					
	485		490		495
Ala Gly Thr Thr Glu Gly Val Thr Ala Leu Gln Met Asp Ile Lys Ile					
	500		505		510
Gln Gly Ile Thr Lys Glu Ile Met Gln Ile Ala Leu Ala Gln Ala Lys					
	515		520		525
Glu Ala Arg Leu His Ile Leu Asp Gln Met Lys Ala Ala Val Ala Gly					
	530		535		540
Pro Gln Glu Leu Ser Ala His Ala Pro Arg Leu Phe Thr Met Lys Ile					
545		550		555	560
Asn Gln Asp Lys Ile Arg Glu Val Ile Gly Lys Gly Gly Glu Thr Ile					
	565		570		575
Arg Ala Ile Thr Ala Glu Thr Gly Thr Glu Ile Asn Ile Ala Glu Asp					

580 585 590
 Gly Thr Ile Thr Ile Ala Ala Thr Thr Gln Glu Ala Gly Asp Ala Ala
 595 600 605
 Lys Lys Arg Ile Glu Gln Ile Thr Ala Glu Val Glu Val Gly Lys Val
 610 615 620
 Tyr Glu Gly Thr Val Val Lys Ile Leu Asp Asn Asn Val Gly Ala Ile
 625 630 635 640
 Val Ser Val Met Pro Gly Lys Asp Gly Leu Val His Ile Ser Gln Ile
 645 650 655
 Ala His Glu Arg Val Arg Asn Val Gly Asp Tyr Leu Gln Val Gly Gln
 660 665 670
 Val Val Asn Val Lys Ala Leu Glu Val Asp Asp Arg Gly Arg Val Arg
 675 680 685
 Leu Ser Ile Lys Ala Leu Leu Asp Ala Pro Ala Arg Glu Glu Asn Ala
 690 695 700
 Ala Glu
 705

<210> 37
 <211> 3972
 <212> DNA
 <213> Neisseria meningitidis

<220>
 <221> CDS
 <222> (1)..(3972)

<400> 37
 gtg agc cga att atg tct gtc gtt ttg ccc ttg cgc gcc gtt acc gcc 48
 Val Ser Arg Ile Met Ser Val Val Leu Pro Leu Arg Gly Val Thr Ala
 1 5 10 15
 ctt tcc gat ttc cgt gtt gaa aaa ctc ttg caa aaa gcc gcc gca ctc 96
 Leu Ser Asp Phe Arg Val Glu Lys Leu Leu Gln Lys Ala Ala Ala Leu
 20 25 30
 ggt ctg ccc gaa gtc aaa tta agc agc gaa ttt tgg tat ttc gtc gcc 144
 Gly Leu Pro Glu Val Lys Leu Ser Ser Glu Phe Trp Tyr Phe Val Gly

35	40	45	
agc gag aaa gca ctt gat gcc gcg act gtc gaa aaa ctg caa gcc ttg			192
Ser Glu Lys Ala Leu Asp Ala Ala Thr Val Glu Lys Leu Gln Ala Leu			
50	55	60	
ttg gcg gcg caa agc gtt gaa caa acg cca aaa gcg cgc gag ggc ttg			240
Leu Ala Ala Gln Ser Val Glu Gln Thr Pro Lys Ala Arg Glu Gly Leu			
65	70	75	80
cat ttg ttt ttg gtc acg ccc cgt ttg ggt acg att tcg ccg tgg gct			288
His Leu Phe Leu Val Thr Pro Arg Leu Gly Thr Ile Ser Pro Trp Ala			
85	90	95	
tcc aag gcg acc aat atc gcg gaa aac tgc ggt ttg gca ggc atc gaa			336
Ser Lys Ala Thr Asn Ile Ala Glu Asn Cys Gly Leu Ala Gly Ile Glu			
100	105	110	
cgc atc gag cgc ggc atg gcg gtg tgg ctg gaa ggt gcg ctt acc gat			384
Arg Ile Glu Arg Gly Met Ala Val Trp Leu Glu Gly Ala Leu Thr Asp			
115	120	125	
gaa cag caa cag caa tgg gcg gct ttg ctg cac gac cgc atg act gaa			432
Glu Gln Gln Gln Gln Trp Ala Ala Leu Leu His Asp Arg Met Thr Glu			
130	135	140	
tcc gtg ctg ccc gat ttt cag acg gca tcc aaa tta ttc cac cat ctc			480
Ser Val Leu Pro Asp Phe Gln Thr Ala Ser Lys Leu Phe His His Leu			
145	150	155	160
gaa tcc gaa acc ttt tcc acc gtc gat gtt ttg ggc ggc ggt aaa gag			528
Glu Ser Glu Thr Phe Ser Thr Val Asp Val Leu Gly Gly Gly Lys Glu			
165	170	175	
gct ttg gtc aaa gcc aat acc gaa atg ggt ttg gca ctc tcc gcc gac			576
Ala Leu Val Lys Ala Asn Thr Glu Met Gly Leu Ala Leu Ser Ala Asp			
180	185	190	
gaa atc gat tat ctg gtt gaa aac tat cag gct ttg cag cgc aat ccg			624
Glu Ile Asp Tyr Leu Val Glu Asn Tyr Gln Ala Leu Gln Arg Asn Pro			
195	200	205	
tct gat gtg gag ctg atg atg ttc gcg cag gca aac agc gaa cac tgc			672
Ser Asp Val Glu Leu Met Met Phe Ala Gln Ala Asn Ser Glu His Cys			
210	215	220	
cgc cac aaa atc ttc aac gcc gac ttc atc ctc aac ggc gaa aaa cag			720
Arg His Lys Ile Phe Asn Ala Asp Phe Ile Leu Asn Gly Glu Lys Gln			

225	230	235	240	
ccg aaa tcg ctc ttc ggt atg ata cgc gac acg cac aac gcg cat ccc				768
Pro Lys Ser Leu Phe Gly Met Ile Arg Asp Thr His Asn Ala His Pro				
245	250	255		
gaa ggc aca gtc gtt gcc tac aaa gac aac tca tcc gtg atc gaa ggc				816
Glu Gly Thr Val Val Ala Tyr Lys Asp Asn Ser Ser Val Ile Glu Gly				
260	265	270		
gca aaa gtc gaa cgt ttc tat cca aat gcg gca gaa aac caa ggc tac				864
Ala Lys Val Glu Arg Phe Tyr Pro Asn Ala Ala Glu Asn Gln Gly Tyr				
275	280	285		
cgt ttc cac gag gaa gac acg cat atc atc atg aaa gtg gaa acg cac				912
Arg Phe His Glu Glu Asp Thr His Ile Ile Met Lys Val Glu Thr His				
290	295	300		
aac cac ccg acc gcc atc gcg ccg ttt gcg ggc gcg gca acg ggc gcg				960
Asn His Pro Thr Ala Ile Ala Pro Phe Ala Gly Ala Ala Thr Gly Ala				
305	310	315	320	
ggc ggc gaa atc cgc gac gaa ggt gcg acg ggc aaa ggt tcg cgt ccg				1008
Gly Gly Glu Ile Arg Asp Glu Gly Ala Thr Gly Lys Gly Ser Arg Pro				
325	330	335		
aaa gca ggc ttg acc ggc ttt acc gtg tcc aac ctc aac atc ccc ggc				1056
Lys Ala Gly Leu Thr Gly Phe Thr Val Ser Asn Leu Asn Ile Pro Gly				
340	345	350		
ctc aaa cag ccg tgg gaa caa gat tac ggc aag ccg gaa cat att tcc				1104
Leu Lys Gln Pro Trp Glu Gln Asp Tyr Gly Lys Pro Glu His Ile Ser				
355	360	365		
tcg ccg ctg gac att atg att gaa ggc ccc atc ggc ggc gcg gcg ttc				1152
Ser Pro Leu Asp Ile Met Ile Glu Gly Pro Ile Gly Gly Ala Ala Phe				
370	375	380		
aac aac gaa ttc ggc cgc ccc aac ctc ttg ggc tac ttc cgc act ttt				1200
Asn Asn Glu Phe Gly Arg Pro Asn Leu Leu Gly Tyr Phe Arg Thr Phe				
385	390	395	400	
gaa gaa aaa ttt gac ggt cag gtt cgc ggc tac cac aaa ccg att atg				1248
Glu Glu Lys Phe Asp Gly Gln Val Arg Gly Tyr His Lys Pro Ile Met				
405	410	415		
att gcc ggc ggc ttg ggc agc att cag gcg cag cag acg cat aaa gac				1296
Ile Ala Gly Gly Leu Gly Ser Ile Gln Ala Gln Gln Thr His Lys Asp				

420	425	430	
gaa atc ccc gaa ggc gcg ttg ctg atc caa ctg ggc ggc ccg ggt atg			1344
Glu Ile Pro Glu Gly Ala Leu Leu Ile Gln Leu Gly Gly Pro Gly Met			
435	440	445	
ctt atc ggc ttg ggc ggc ggc gcg gct tct tcg atg gat acc ggc aca			1392
Leu Ile Gly Leu Gly Gly Gly Ala Ala Ser Ser Met Asp Thr Gly Thr			
450	455	460	
aac gac gcg tct tta gac ttc aac tcc gta caa cgc ggc aac ccc gaa			1440
Asn Asp Ala Ser Leu Asp Phe Asn Ser Val Gln Arg Gly Asn Pro Glu			
465	470	475	480
atc gaa cgc cgc gcg cag gaa gtg atc gac cgc tgc tgg cag ctc ggc			1488
Ile Glu Arg Arg Ala Gln Glu Val Ile Asp Arg Cys Trp Gln Leu Gly			
485	490	495	
gac aaa aac ccg att atc tcc atc cac gac gtt ggc gcg ggc ggc ttg			1536
Asp Lys Asn Pro Ile Ile Ser Ile His Asp Val Gly Ala Gly Gly Leu			
500	505	510	
tcc aac gcc ttc ccc gaa ctc gtc aac gat gcc gga cgc ggc gcg gta			1584
Ser Asn Ala Phe Pro Glu Leu Val Asn Asp Ala Gly Arg Gly Ala Val			
515	520	525	
ttc aag ctg cgc gaa gtg ccg ctg gaa gaa cac ggt ctc aac ccg ctg			1632
Phe Lys Leu Arg Glu Val Pro Leu Glu Glu His Gly Leu Asn Pro Leu			
530	535	540	
caa atc tgg tgc aac gaa tcg caa gag cgt tat gta ttg tcg att ttg			1680
Gln Ile Trp Cys Asn Glu Ser Gln Glu Arg Tyr Val Leu Ser Ile Leu			
545	550	555	560
gaa aaa gat ttg gac atc ttc cgc tcc atc tgc gaa cgt gaa cgc tgc			1728
Glu Lys Asp Leu Asp Ile Phe Arg Ser Ile Cys Glu Arg Glu Arg Cys			
565	570	575	
ccg ttt gcg gta gtc ggt acg gcg acc gac gac ggc cat ttg aaa gta			1776
Pro Phe Ala Val Val Gly Thr Ala Thr Asp Asp Gly His Leu Lys Val			
580	585	590	
cgc gac gat ttg ttc tcc aac aac cct gtc gat ttg ccg ttg aac gtc			1824
Arg Asp Asp Leu Phe Ser Asn Asn Pro Val Asp Leu Pro Leu Asn Val			
595	600	605	
ttg ctc ggc aaa ccg ccc aaa acc acg cgt acc gac aaa acg gtt gcg			1872
Leu Leu Gly Lys Pro Pro Lys Thr Thr Arg Thr Asp Lys Thr Val Ala			

610	615	620	
ccg tcc aaa aaa ccg ttt cac gcg ggc gat atc gac att act gaa gcc			1920
Pro Ser Lys Lys Pro Phe His Ala Gly Asp Ile Asp Ile Thr Glu Ala			
625	630	635	640
gcc tac cgc gtt ctg cgc ctg cct gcc gta gcc gcc aaa aac ttc ctg			1968
Ala Tyr Arg Val Leu Arg Leu Pro Ala Val Ala Ala Lys Asn Phe Leu			
645	650	655	
att acc atc ggc gac cgc agc gtc ggc ggc atg acc cac cgc gac caa			2016
Ile Thr Ile Gly Asp Arg Ser Val Gly Gly Met Thr His Arg Asp Gln			
660	665	670	
atg gtc ggc aaa tac caa acc ccc gta gcc gac tgc gcc gtt acc atg			2064
Met Val Gly Lys Tyr Gln Thr Pro Val Ala Asp Cys Ala Val Thr Met			
675	680	685	
atg ggc ttc aac act tat cgc ggc gaa gcg atg agc atg ggc gaa aaa			2112
Met Gly Phe Asn Thr Tyr Arg Gly Glu Ala Met Ser Met Gly Glu Lys			
690	695	700	
ccg acc gtc gcc ctg ttt gac gcg cct gct tcg ggc aga atg tgc gtc			2160
Pro Thr Val Ala Leu Phe Asp Ala Pro Ala Ser Gly Arg Met Cys Val			
705	710	715	720
ggc gaa gcc atc acc aac atc gcg gca gcc aat atc ggc gac atc ggc			2208
Gly Glu Ala Ile Thr Asn Ile Ala Ala Ala Asn Ile Gly Asp Ile Gly			
725	730	735	
aac atc aaa ctc tcc gcc aac tgg atg gcg gcg tgc ggc aac gaa ggc			2256
Asn Ile Lys Leu Ser Ala Asn Trp Met Ala Ala Cys Gly Asn Glu Gly			
740	745	750	
gaa gac gaa aaa ctc tac cgc acc gtc gaa gcg gtt tcc aaa gcc tgt			2304
Glu Asp Glu Lys Leu Tyr Arg Thr Val Glu Ala Val Ser Lys Ala Cys			
755	760	765	
cag gca ttg gat ttg agc att ccc gtg ggc aaa gac agc ctg tcg atg			2352
Gln Ala Leu Asp Leu Ser Ile Pro Val Gly Lys Asp Ser Leu Ser Met			
770	775	780	
aaa acc gtg tgg cag gac ggc gaa gag aaa aaa tcc gtc gtt tcg ccg			2400
Lys Thr Val Trp Gln Asp Gly Glu Glu Lys Lys Ser Val Val Ser Pro			
785	790	795	800
ttg agc ctg att atc tcc gcg ttc gcg ccg gtt caa gac gta cgc aaa			2448
Leu Ser Leu Ile Ile Ser Ala Phe Ala Pro Val Gln Asp Val Arg Lys			

	805	810	815	
acc gtt acg ccc gaa ttg aaa aac gtc gag gac agc gta ttg ctg ttt				2496
Thr Val Thr Pro Glu Leu Lys Asn Val Glu Asp Ser Val Leu Leu Phe				
	820	825	830	
gtc gat ttg ggc ttc ggc aaa gcg cgt atg ggc ggc tcg gcg ttt ggt				2544
Val Asp Leu Gly Phe Gly Lys Ala Arg Met Gly Gly Ser Ala Phe Gly				
	835	840	845	
cag gtg tac aac aat atg agc ggc gac gcg ccc gat ttg gac gat aca				2592
Gln Val Tyr Asn Asn Met Ser Gly Asp Ala Pro Asp Leu Asp Asp Thr				
	850	855	860	
agt cgt ctg aaa gcg ttt tac aac gtg att cag cag ctt gtc gcc gaa				2640
Ser Arg Leu Lys Ala Phe Tyr Asn Val Ile Gln Gln Leu Val Ala Glu				
	865	870	875	880
gac aaa ctc ttg gcg tat cac gac cgc agc gac ggc ggc ttg ttt gcc				2688
Asp Lys Leu Leu Ala Tyr His Asp Arg Ser Asp Gly Gly Leu Phe Ala				
	885	890	895	
gtt ttg gta gaa atg gcg ttt gca ggg cgg tgc ggc ttg gat ata gat				2736
Val Leu Val Glu Met Ala Phe Ala Gly Arg Cys Gly Leu Asp Ile Asp				
	900	905	910	
tta aat tta ttg ctt gca caa aca ttt att acc aac cat acc gct ctg				2784
Leu Asn Leu Leu Leu Ala Gln Thr Phe Ile Thr Asn His Thr Ala Leu				
	915	920	925	
tct caa tca ttg cgg act gaa gag gta aaa gcg ttg gct gaa tgg caa				2832
Ser Gln Ser Leu Arg Thr Glu Glu Val Lys Ala Leu Ala Glu Trp Gln				
	930	935	940	
gaa acc att gcc cgc aca tta ttt aat gaa gag ttg ggt gct gtt atc				2880
Glu Thr Ile Ala Arg Thr Leu Phe Asn Glu Glu Leu Gly Ala Val Ile				
	945	950	955	960
caa gtt aga aaa caa gat gtt gcc gat att atc aat tta ttc tat caa				2928
Gln Val Arg Lys Gln Asp Val Ala Asp Ile Ile Asn Leu Phe Tyr Gln				
	965	970	975	
caa cag ctg cat cat aat gtc ttt gaa atc ggt acg tta act gat gag				2976
Gln Gln Leu His His Asn Val Phe Glu Ile Gly Thr Leu Thr Asp Glu				
	980	985	990	
aac acg tta atc atc cgc gac ggg caa acg cac ctt att tct gac aac				3024
Asn Thr Leu Ile Ile Arg Asp Gly Gln Thr His Leu Ile Ser Asp Asn				

995	1000	1005	
cta atc aaa ctg caa caa acc tgg caa gaa acc agc cat caa atc caa			3072
Leu Ile Lys Leu Gln Gln Thr Trp Gln Glu Thr Ser His Gln Ile Gln			
1010	1015	1020	
cgc ctg cgc gac aac cct gcc tgc gcc gac agc gag ttt gcc ctg att			3120
Arg Leu Arg Asp Asn Pro Ala Cys Ala Asp Ser Glu Phe Ala Leu Ile			
1025	1030	1035	1040
ggc gac aac gga cgc agc gca ttg ttt gcc aac ctg aaa ttc gac gtg			3168
Gly Asp Asn Gly Arg Ser Ala Leu Phe Ala Asn Leu Lys Phe Asp Val			
1045	1050	1055	
aac gaa gac atc gcc gcg cgc ttt atc aac agc ggc gcg aaa ccc aaa			3216
Asn Glu Asp Ile Ala Ala Pro Phe Ile Asn Ser Gly Ala Lys Pro Lys			
1060	1065	1070	
atc gcc atc ctg cgc gaa cag ggc gta aac ggg caa atc gaa atg gcc			3264
Ile Ala Ile Leu Arg Glu Gln Gly Val Asn Gly Gln Ile Glu Met Ala			
1075	1080	1085	
gcc gcc ttc acc cgt gcc ggt ttc gat gcc tac gac gtg cat atg tcc			3312
Ala Ala Phe Thr Arg Ala Gly Phe Asp Ala Tyr Asp Val His Met Ser			
1090	1095	1100	
gac ctg atg gca ggc cgc gtc cac ctt gcc gac ttc aaa atg ctg gcg			3360
Asp Leu Met Ala Gly Arg Val His Leu Ala Asp Phe Lys Met Leu Ala			
1105	1110	1115	1120
gcg tgc ggc ggc ttc agc tac ggc gac gta ctc ggc gcg ggc aaa ggc			3408
Ala Cys Gly Gly Phe Ser Tyr Gly Asp Val Leu Gly Ala Gly Lys Gly			
1125	1130	1135	
tgg gcg aaa tcc atc ctg ttc cac ccc gcc tta cgc gac cag ttc gcc			3456
Trp Ala Lys Ser Ile Leu Phe His Pro Ala Leu Arg Asp Gln Phe Ala			
1140	1145	1150	
gcc ttc ttc acc gac ccg aac acg ctg aca ttg ggc gtg tgc aac ggc			3504
Ala Phe Phe Thr Asp Pro Asn Thr Leu Thr Leu Gly Val Cys Asn Gly			
1155	1160	1165	
tgc cag atg gtc agc aac ctc gcc gaa atc atc ccc ggc acg gca ggc			3552
Cys Gln Met Val Ser Asn Leu Ala Glu Ile Ile Pro Gly Thr Ala Gly			
1170	1175	1180	
tgg ccg aag ttc aag cgc aac ctg agc gaa cag ttc gaa gcg cgc ctg			3600
Trp Pro Lys Phe Lys Arg Asn Leu Ser Glu Gln Phe Glu Ala Arg Leu			

1185	1190	1195	1200	
agc atg gtt cac gtc ccc aaa tca gcc tcc ctg att ctg aac gaa atg				3648
Ser Met Val His Val Pro Lys Ser Ala Ser Leu Ile Leu Asn Glu Met				
1205		1210	1215	
caa ggc tcc agc ctg ccc gtc gtc gtc agc cac ggc gaa ggc cgc gcc				3696
Gln Gly Ser Ser Leu Pro Val Val Val Ser His Gly Glu Gly Arg Ala				
1220		1225	1230	
gac ttc gcg ctt cac ggc ggc aac att tct gcc gat ttg ggc att gcg				3744
Asp Phe Ala Leu His Gly Gly Asn Ile Ser Ala Asp Leu Gly Ile Ala				
1235		1240	1245	
ttg caa tat gtg gac gga caa aac caa att acc caa acc tac ccg ctc				3792
Leu Gln Tyr Val Asp Gly Gln Asn Gln Ile Thr Gln Thr Tyr Pro Leu				
1250		1255	1260	
aac ccc aac ggc tcg ccg caa ggc atc gcc ggc gtg acc aac gcc gac				3840
Asn Pro Asn Gly Ser Pro Gln Gly Ile Ala Gly Val Thr Asn Ala Asp				
1265		1270	1275	1280
ggc cgc gtt acc atc atg atg ccg cat cct gaa cgt gta tac cgc gcc				3888
Gly Arg Val Thr Ile Met Met Pro His Pro Glu Arg Val Tyr Arg Ala				
1285		1290	1295	
gca caa atg agc tgg aaa ccg gaa gac tgg acg gaa ttg tcc ggc tgg				3936
Ala Gln Met Ser Trp Lys Pro Glu Asp Trp Thr Glu Leu Ser Gly Trp				
1300		1305	1310	
tac cgc ctc ttc gcc ggc gca cgt aaa gcc ttg ggc				3972
Tyr Arg Leu Phe Ala Gly Ala Arg Lys Ala Leu Gly				
1315		1320		

<210> 38

<211> 1324

<212> PRT

<213> Neisseria meningitidis

<400> 38

Val Ser Arg Ile Met Ser Val Val Leu Pro Leu Arg Gly Val Thr Ala
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Leu Ser Asp Phe Arg Val Glu Lys Leu Leu Gln Lys Ala Ala Ala Leu
20 25 30

Gly Leu Pro Glu Val Lys Leu Ser Ser Glu Phe Trp Tyr Phe Val Gly

35	40	45
Ser Glu Lys Ala Leu Asp Ala Ala Thr Val Glu Lys Leu Gln Ala Leu		
50	55	60
Leu Ala Ala Gln Ser Val Glu Gln Thr Pro Lys Ala Arg Glu Gly Leu		
65	70	75 80
His Leu Phe Leu Val Thr Pro Arg Leu Gly Thr Ile Ser Pro Trp Ala		
85	90	95
Ser Lys Ala Thr Asn Ile Ala Glu Asn Cys Gly Leu Ala Gly Ile Glu		
100	105	110
Arg Ile Glu Arg Gly Met Ala Val Trp Leu Glu Gly Ala Leu Thr Asp		
115	120	125
Glu Gln Gln Gln Gln Trp Ala Ala Leu Leu His Asp Arg Met Thr Glu		
130	135	140
Ser Val Leu Pro Asp Phe Gln Thr Ala Ser Lys Leu Phe His His Leu		
145	150	155 160
Glu Ser Glu Thr Phe Ser Thr Val Asp Val Leu Gly Gly Gly Lys Glu		
165	170	175
Ala Leu Val Lys Ala Asn Thr Glu Met Gly Leu Ala Leu Ser Ala Asp		
180	185	190
Glu Ile Asp Tyr Leu Val Glu Asn Tyr Gln Ala Leu Gln Arg Asn Pro		
195	200	205
Ser Asp Val Glu Leu Met Met Phe Ala Gln Ala Asn Ser Glu His Cys		
210	215	220
Arg His Lys Ile Phe Asn Ala Asp Phe Ile Leu Asn Gly Glu Lys Gln		
225	230	235 240
Pro Lys Ser Leu Phe Gly Met Ile Arg Asp Thr His Asn Ala His Pro		
245	250	255
Glu Gly Thr Val Val Ala Tyr Lys Asp Asn Ser Ser Val Ile Glu Gly		
260	265	270
Ala Lys Val Glu Arg Phe Tyr Pro Asn Ala Ala Glu Asn Gln Gly Tyr		
275	280	285
Arg Phe His Glu Glu Asp Thr His Ile Ile Met Lys Val Glu Thr His		

290	295	300
Asn His Pro Thr Ala Ile Ala Pro Phe Ala Gly Ala Ala Thr Gly Ala		
305	310	315 320
Gly Gly Glu Ile Arg Asp Glu Gly Ala Thr Gly Lys Gly Ser Arg Pro		
	325	330 335
Lys Ala Gly Leu Thr Gly Phe Thr Val Ser Asn Leu Asn Ile Pro Gly		
	340	345 350
Leu Lys Gln Pro Trp Glu Gln Asp Tyr Gly Lys Pro Glu His Ile Ser		
	355	360 365
Ser Pro Leu Asp Ile Met Ile Glu Gly Pro Ile Gly Gly Ala Ala Phe		
	370	375 380
Asn Asn Glu Phe Gly Arg Pro Asn Leu Leu Gly Tyr Phe Arg Thr Phe		
385	390	395 400
Glu Glu Lys Phe Asp Gly Gln Val Arg Gly Tyr His Lys Pro Ile Met		
	405	410 415
Ile Ala Gly Gly Leu Gly Ser Ile Gln Ala Gln Gln Thr His Lys Asp		
	420	425 430
Glu Ile Pro Glu Gly Ala Leu Leu Ile Gln Leu Gly Gly Pro Gly Met		
	435	440 445
Leu Ile Gly Leu Gly Gly Gly Ala Ala Ser Ser Met Asp Thr Gly Thr		
	450	455 460
Asn Asp Ala Ser Leu Asp Phe Asn Ser Val Gln Arg Gly Asn Pro Glu		
465	470	475 480
Ile Glu Arg Arg Ala Gln Glu Val Ile Asp Arg Cys Trp Gln Leu Gly		
	485	490 495
Asp Lys Asn Pro Ile Ile Ser Ile His Asp Val Gly Ala Gly Gly Leu		
	500	505 510
Ser Asn Ala Phe Pro Glu Leu Val Asn Asp Ala Gly Arg Gly Ala Val		
	515	520 525
Phe Lys Leu Arg Glu Val Pro Leu Glu Glu His Gly Leu Asn Pro Leu		
	530	535 540
Gln Ile Trp Cys Asn Glu Ser Gln Glu Arg Tyr Val Leu Ser Ile Leu		

545	550	555	560
Glu Lys Asp Leu Asp Ile Phe Arg Ser Ile Cys Glu Arg Glu Arg Cys			
	565	570	575
Pro Phe Ala Val Val Gly Thr Ala Thr Asp Asp Gly His Leu Lys Val			
	580	585	590
Arg Asp Asp Leu Phe Ser Asn Asn Pro Val Asp Leu Pro Leu Asn Val			
	595	600	605
Leu Leu Gly Lys Pro Pro Lys Thr Thr Arg Thr Asp Lys Thr Val Ala			
	610	615	620
Pro Ser Lys Lys Pro Phe His Ala Gly Asp Ile Asp Ile Thr Glu Ala			
	625	630	635
Ala Tyr Arg Val Leu Arg Leu Pro Ala Val Ala Ala Lys Asn Phe Leu			
	645	650	655
Ile Thr Ile Gly Asp Arg Ser Val Gly Gly Met Thr His Arg Asp Gln			
	660	665	670
Met Val Gly Lys Tyr Gln Thr Pro Val Ala Asp Cys Ala Val Thr Met			
	675	680	685
Met Gly Phe Asn Thr Tyr Arg Gly Glu Ala Met Ser Met Gly Glu Lys			
	690	695	700
Pro Thr Val Ala Leu Phe Asp Ala Pro Ala Ser Gly Arg Met Cys Val			
	705	710	715
Gly Glu Ala Ile Thr Asn Ile Ala Ala Ala Asn Ile Gly Asp Ile Gly			
	725	730	735
Asn Ile Lys Leu Ser Ala Asn Trp Met Ala Ala Cys Gly Asn Glu Gly			
	740	745	750
Glu Asp Glu Lys Leu Tyr Arg Thr Val Glu Ala Val Ser Lys Ala Cys			
	755	760	765
Gln Ala Leu Asp Leu Ser Ile Pro Val Gly Lys Asp Ser Leu Ser Met			
	770	775	780
Lys Thr Val Trp Gln Asp Gly Glu Glu Lys Lys Ser Val Val Ser Pro			
	785	790	795
Leu Ser Leu Ile Ile Ser Ala Phe Ala Pro Val Gln Asp Val Arg Lys			

	805		810		815
Thr Val Thr Pro Glu Leu Lys Asn Val Glu Asp Ser Val Leu Leu Phe	820		825		830
Val Asp Leu Gly Phe Gly Lys Ala Arg Met Gly Gly Ser Ala Phe Gly	835		840		845
Gln Val Tyr Asn Asn Met Ser Gly Asp Ala Pro Asp Leu Asp Asp Thr	850		855		860
Ser Arg Leu Lys Ala Phe Tyr Asn Val Ile Gln Gln Leu Val Ala Glu	865		870		875
					880
Asp Lys Leu Leu Ala Tyr His Asp Arg Ser Asp Gly Gly Leu Phe Ala	885		890		895
Val Leu Val Glu Met Ala Phe Ala Gly Arg Cys Gly Leu Asp Ile Asp	900		905		910
Leu Asn Leu Leu Leu Ala Gln Thr Phe Ile Thr Asn His Thr Ala Leu	915		920		925
Ser Gln Ser Leu Arg Thr Glu Glu Val Lys Ala Leu Ala Glu Trp Gln	930		935		940
Glu Thr Ile Ala Arg Thr Leu Phe Asn Glu Glu Leu Gly Ala Val Ile	945		950		955
					960
Gln Val Arg Lys Gln Asp Val Ala Asp Ile Ile Asn Leu Phe Tyr Gln	965		970		975
Gln Gln Leu His His Asn Val Phe Glu Ile Gly Thr Leu Thr Asp Glu	980		985		990
Asn Thr Leu Ile Ile Arg Asp Gly Gln Thr His Leu Ile Ser Asp Asn	995		1000		1005
Leu Ile Lys Leu Gln Gln Thr Trp Gln Glu Thr Ser His Gln Ile Gln	1010		1015		1020
Arg Leu Arg Asp Asn Pro Ala Cys Ala Asp Ser Glu Phe Ala Leu Ile	1025		1030		1035
					1040
Gly Asp Asn Gly Arg Ser Ala Leu Phe Ala Asn Leu Lys Phe Asp Val	1045		1050		1055
Asn Glu Asp Ile Ala Ala Pro Phe Ile Asn Ser Gly Ala Lys Pro Lys					

1060	1065	1070
Ile Ala Ile Leu Arg Glu Gln Gly Val Asn Gly Gln Ile Glu Met Ala		
1075	1080	1085
Ala Ala Phe Thr Arg Ala Gly Phe Asp Ala Tyr Asp Val His Met Ser		
1090	1095	1100
Asp Leu Met Ala Gly Arg Val His Leu Ala Asp Phe Lys Met Leu Ala		
1105	1110	1115
Ala Cys Gly Gly Phe Ser Tyr Gly Asp Val Leu Gly Ala Gly Lys Gly		
1125	1130	1135
Trp Ala Lys Ser Ile Leu Phe His Pro Ala Leu Arg Asp Gln Phe Ala		
1140	1145	1150
Ala Phe Phe Thr Asp Pro Asn Thr Leu Thr Leu Gly Val Cys Asn Gly		
1155	1160	1165
Cys Gln Met Val Ser Asn Leu Ala Glu Ile Ile Pro Gly Thr Ala Gly		
1170	1175	1180
Trp Pro Lys Phe Lys Arg Asn Leu Ser Glu Gln Phe Glu Ala Arg Leu		
1185	1190	1195
Ser Met Val His Val Pro Lys Ser Ala Ser Leu Ile Leu Asn Glu Met		
1205	1210	1215
Gln Gly Ser Ser Leu Pro Val Val Val Ser His Gly Glu Gly Arg Ala		
1220	1225	1230
Asp Phe Ala Leu His Gly Gly Asn Ile Ser Ala Asp Leu Gly Ile Ala		
1235	1240	1245
Leu Gln Tyr Val Asp Gly Gln Asn Gln Ile Thr Gln Thr Tyr Pro Leu		
1250	1255	1260
Asn Pro Asn Gly Ser Pro Gln Gly Ile Ala Gly Val Thr Asn Ala Asp		
1265	1270	1275
Gly Arg Val Thr Ile Met Met Pro His Pro Glu Arg Val Tyr Arg Ala		
1285	1290	1295
Ala Gln Met Ser Trp Lys Pro Glu Asp Trp Thr Glu Leu Ser Gly Trp		
1300	1305	1310
Tyr Arg Leu Phe Ala Gly Ala Arg Lys Ala Leu Gly		

1315

1320

<210> 39

<211> 471

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(471)

<400> 39

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Met Ser Arg Asn Thr Glu Glu Leu Gln Gly Ile Ser Leu Leu Gly Asn

1

5

10

15

cag aaa acc cga tat ccg acc ggc tac gcg cct gaa att ctc gaa gca 96

Gln Lys Thr Arg Tyr Pro Thr Gly Tyr Ala Pro Glu Ile Leu Glu Ala

20

25

30

ttc gac aac aaa cat ccc gac aac gac tat ttc gtc aaa ttc gtc tgc 144

Phe Asp Asn Lys His Pro Asp Asn Asp Tyr Phe Val Lys Phe Val Cys

35

40

45

cca gag ttc acc agc ctc tgc ccc atg acc ggg cag ccc gac ttc gcc 192

Pro Glu Phe Thr Ser Leu Cys Pro Met Thr Gly-Gln Pro Asp Phe Ala

50

55

60

acc atc tac atc cgc tac att ccg cac atc aaa atg gtg gaa agc aaa 240

Thr Ile Tyr Ile Arg Tyr Ile Pro His Ile Lys Met Val Glu Ser Lys

65

70

75

80

tcc ctg aaa ctc tac ctc ttc agc ttc cgc aac cac ggc gat ttt cat 288

Ser Leu Lys Leu Tyr Leu Phe Ser Phe Arg Asn His Gly Asp Phe His

85

90

95

gaa gac tgc gtc aac atc atc atg aaa gac ctc att gcc ctg atg gat 336

Glu Asp Cys Val Asn Ile Ile Met Lys Asp Leu Ile Ala Leu Met Asp

100

105

110

ccg aaa tac atc gaa gta ttc ggc gag ttc aca ccg cgc ggc ggc atc 384

Pro Lys Tyr Ile Glu Val Phe Gly Glu Phe Thr Pro Arg Gly Gly Ile

115

120

125

gcc att cat cct ttc gcc aat tac ggc aaa gca ggc acc gag ttt gaa 432

Ala Ile His Pro Phe Ala Asn Tyr Gly Lys Ala Gly Thr Glu Phe Glu

130 135 140 471

gca ttg gcg cgt aaa cgc ctg ttt gag cac gac gca caa
 Ala Leu Ala Arg Lys Arg Leu Phe Glu His Asp Ala Gln
 145 150 155

<210> 40
 <211> 157
 <212> PRT
 <213> Neisseria meningitidis

<400> 40
 Met Ser Arg Asn Thr Glu Glu Leu Gln Gly Ile Ser Leu Leu Gly Asn
 1 5 10 15

Gln Lys Thr Arg Tyr Pro Thr Gly Tyr Ala Pro Glu Ile Leu Glu Ala
 20 25 30

Phe Asp Asn Lys His Pro Asp Asn Asp Tyr Phe Val Lys Phe Val Cys
 35 40 45

Pro Glu Phe Thr Ser Leu Cys Pro Met Thr Gly Gln Pro Asp Phe Ala
 50 55 60

Thr Ile Tyr Ile Arg Tyr Ile Pro His Ile Lys Met Val Glu Ser Lys
 65 70 75 80

Ser Leu Lys Leu Tyr Leu Phe Ser Phe Arg Asn His Gly Asp Phe His
 85 90 95

Glu Asp Cys Val Asn Ile Ile Met Lys Asp Leu Ile Ala Leu Met Asp
 100 105 110

Pro Lys Tyr Ile Glu Val Phe Gly Glu Phe Thr Pro Arg Gly Gly Ile
 115 120 125

Ala Ile His Pro Phe Ala Asn Tyr Gly Lys Ala Gly Thr Glu Phe Glu
 130 135 140

Ala Leu Ala Arg Lys Arg Leu Phe Glu His Asp Ala Gln
 145 150 155

<210> 41
 <211> 504
 <212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(504)

<400> 41

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 1 5 10 15

ccc gcc gta cgc gtg gcg aaa acc atg acc acg ccc aaa ggc gac acc 96
 Pro Ala Val Arg Val Ala Lys Thr Met Thr Thr Pro Lys Gly Asp Thr
 20 25 30

att acc gtg ttc gac ctg cgc ttt tgc gtt ccc aat aaa gaa atc ctg 144
 Ile Thr Val Phe Asp Leu Arg Phe Cys Val Pro Asn Lys Glu Ile Leu
 35 40 45

cct gaa aaa ggc ata cac acg ctg gag cat ttg ttc gca ggt ttt atg 192
 Pro Glu Lys Gly Ile His Thr Leu Glu His Leu Phe Ala Gly Phe Met
 50 55 60

cgc gac cac ttg aac ggc aac ggc gtg gaa atc atc gac att tcc ccg 240
 Arg Asp His Leu Asn Gly Asn Gly Val Glu Ile Ile Asp Ile Ser Pro
 65 70 75 80

atg ggc tgc cgc acc ggt ttt tat atg agc ctt atc ggc acg cct tcc 288
 Met Gly Cys Arg Thr Gly Phe Tyr Met Ser Leu Ile Gly Thr Pro Ser
 85 90 95

gaa cag cag gtc gcc gat gcg tgg ctg gct tcg atg cag gat gtt ttg 336
 Glu Gln Gln Val Ala Asp Ala Trp Leu Ala Ser Met Gln Asp Val Leu
 100 105 110

aat gtc aaa gac caa agc aaa atc ccc gag ttg aac gaa tac caa tgc 384
 Asn Val Lys Asp Gln Ser Lys Ile Pro Glu Leu Asn Glu Tyr Gln Cys
 115 120 125

ggc act tat caa atg cac tcg ctc gcc gaa gcg cag caa atc gcg caa 432
 Gly Thr Tyr Gln Met His Ser Leu Ala Glu Ala Gln Gln Ile Ala Gln
 130 135 140

aac gtg ttg gcg cgc aaa gtg gcg gtg aac aaa aat gaa gag ctg acg 480
 Asn Val Leu Ala Arg Lys Val Ala Val Asn Lys Asn Glu Glu Leu Thr
 145 150 155 160

ctg gat gaa ggg ctg ctg aac gcc 504

Leu Asp Glu Gly Leu Leu Asn Ala
165

<210> 42

<211> 168

<212> PRT

<213> Neisseria meningitidis

<400> 42

Met Pro Leu Leu Asp Ser Phe Lys Val Asp His Thr Arg Met His Ala
1 5 10 15

Pro Ala Val Arg Val Ala Lys Thr Met Thr Thr Pro Lys Gly Asp Thr
20 25 30

Ile Thr Val Phe Asp Leu Arg Phe Cys Val Pro Asn Lys Glu Ile Leu
35 40 45

Pro Glu Lys Gly Ile His Thr Leu Glu His Leu Phe Ala Gly Phe Met
50 55 60

Arg Asp His Leu Asn Gly Asn Gly Val Glu Ile Ile Asp Ile Ser Pro
65 70 75 80

Met Gly Cys Arg Thr Gly Phe Tyr Met Ser Leu Ile Gly Thr Pro Ser
85 90 95

Glu Gln Gln Val Ala Asp Ala Trp Leu Ala Ser Met Gln Asp Val Leu
100 105 110

Asn Val Lys Asp Gln Ser Lys Ile Pro Glu Leu Asn Glu Tyr Gln Cys
115 120 125

Gly Thr Tyr Gln Met His Ser Leu Ala Glu Ala Gln Gln Ile Ala Gln
130 135 140

Asn Val Leu Ala Arg Lys Val Ala Val Asn Lys Asn Glu Glu Leu Thr
145 150 155 160

Leu Asp Glu Gly Leu Leu Asn Ala
165

<210> 43

<211> 813

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(813)

<400> 43

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atg act atg cac gcc ctc ccc cgc tac gcc gtt ttt ggc aac ccc gtc 48
Met Thr Met His Ala Leu Pro Arg Tyr Ala Val Phe Gly Asn Pro Val
  1             5             10             15

gcc cac agc aaa tcg ccg caa att cat caa caa ttt gcc ctt cag gaa 96
Ala His Ser Lys Ser Pro Gln Ile His Gln Gln Phe Ala Leu Gln Glu
      20             25             30

ggc gtt gac att gaa tac gaa cgc att tgc gcc gac atc ggc ggt ttc 144
Gly Val Asp Ile Glu Tyr Glu Arg Ile Cys Ala Asp Ile Gly Gly Phe
      35             40             45

gcg cag gcg gtt tcg aca ttt ttt gaa aca gcc ggt tgc ggg gca aac 192
Ala Gln Ala Val Ser Thr Phe Phe Glu Thr Gly Gly Cys Gly Ala Asn
      50             55             60

gtt acc gta ccg ttc aag cag gaa gcg ttt cat ctg gcg gac gag cat 240
Val Thr Val Pro Phe Lys Gln Glu Ala Phe His Leu Ala Asp Glu His
      65             70             75             80

tct gaa cgc gca ttg gct gcc ggc gcg gtc aac acg ctg att ttt ctg 288
Ser Glu Arg Ala Leu Ala Ala Gly Ala Val Asn Thr Leu Ile Phe Leu
      85             90             95

aaa aac gga aaa ctg cgc gcc ggc gac aat acc gac ggt atc ggt ttg gcc 336
Lys Asn Gly Lys Leu Arg Gly Asp Asn Thr Asp Gly Ile Gly Leu Ala
      100             105             110

aac gac atc acg cag gtc aaa aac att gcc atc gaa gcc aaa acc atc 384
Asn Asp Ile Thr Gln Val Lys Asn Ile Ala Ile Glu Gly Lys Thr Ile
      115             120             125

ttg ctt ttg ggc gcg gcc gcc gcc gcg gtg cgc gcc gtg att cct gtt ttg 432
Leu Leu Leu Gly Ala Gly Gly Ala Val Arg Gly Val Ile Pro Val Leu
      130             135             140

aaa gaa cac cgt cct gcc cgt atc gtc att gcc aac cgt acc cgc gcc 480
Lys Glu His Arg Pro Ala Arg Ile Val Ile Ala Asn Arg Thr Arg Ala
      145             150             155             160

aaa gcc gag gaa ttg gcg cag ctt ttc gcc att gaa gcc gtc ccg atg 528

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Lys Ala Glu Glu Leu Ala Gln Leu Phe Gly Ile Glu Ala Val Pro Met
 165 170 175

gcg gat gtg aac ggc ggt ttt gat atc atc atc aac ggc acg tcg ggc 576
 Ala Asp Val Asn Gly Gly Phe Asp Ile Ile Ile Asn Gly Thr Ser Gly
 180 185 190

ggt cta aac ggt cag att ccc gat att ccg ccc gat att ttt caa aac 624
 Gly Leu Asn Gly Gln Ile Pro Asp Ile Pro Pro Asp Ile Phe Gln Asn
 195 200 205

tgc gcg ctt gcc tac gat atg gtg tac ggc tgc gcg gca aaa ccg ttt 672
 Cys Ala Leu Ala Tyr Asp Met Val Tyr Gly Cys Ala Ala Lys Pro Phe
 210 215 220

tta gat ttt gca cga caa tcg ggt gcg aaa aaa act gcc gac gga ctg 720
 Leu Asp Phe Ala Arg Gln Ser Gly Ala Lys Lys Thr Ala Asp Gly Leu
 225 230 235 240

ggt atg cta gtc ggt caa gcg gcg gct tcc tac gcc ctc tgg cgc gga 768
 Gly Met Leu Val Gly Gln Ala Ala Ala Ser Tyr Ala Leu Trp Arg Gly
 245 250 255

ttt acg ccc gat atc cgc ccc gtt atc gaa tac atg aaa gcc cta 813
 Phe Thr Pro Asp Ile Arg Pro Val Ile Glu Tyr Met Lys Ala Leu
 260 265 270

<210> 44
 <211> 271
 <212> PRT
 <213> Neisseria meningitidis

<400> 44
 Met Thr Met His Ala Leu Pro Arg Tyr Ala Val Phe Gly Asn Pro Val
 1 5 10 15

Ala His Ser Lys Ser Pro Gln Ile His Gln Gln Phe Ala Leu Gln Glu
 20 25 30

Gly Val Asp Ile Glu Tyr Glu Arg Ile Cys Ala Asp Ile Gly Gly Phe
 35 40 45

Ala Gln Ala Val Ser Thr Phe Phe Glu Thr Gly Gly Cys Gly Ala Asn
 50 55 60

Val Thr Val Pro Phe Lys Gln Glu Ala Phe His Leu Ala Asp Glu His
 65 70 75 80

Ser Glu Arg Ala Leu Ala Ala Gly Ala Val Asn Thr Leu Ile Phe Leu
 85 90 95
 Lys Asn Gly Lys Leu Arg Gly Asp Asn Thr Asp Gly Ile Gly Leu Ala
 100 105 110
 Asn Asp Ile Thr Gln Val Lys Asn Ile Ala Ile Glu Gly Lys Thr Ile
 115 120 125
 Leu Leu Leu Gly Ala Gly Gly Ala Val Arg Gly Val Ile Pro Val Leu
 130 135 140
 Lys Glu His Arg Pro Ala Arg Ile Val Ile Ala Asn Arg Thr Arg Ala
 145 150 155 160
 Lys Ala Glu Glu Leu Ala Gln Leu Phe Gly Ile Glu Ala Val Pro Met
 165 170 175
 Ala Asp Val Asn Gly Gly Phe Asp Ile Ile Ile Asn Gly Thr Ser Gly
 180 185 190
 Gly Leu Asn Gly Gln Ile Pro Asp Ile Pro Pro Asp Ile Phe Gln Asn
 195 200 205
 Cys Ala Leu Ala Tyr Asp Met Val Tyr Gly Cys Ala Ala Lys Pro Phe
 210 215 220
 Leu Asp Phe Ala Arg Gln Ser Gly Ala Lys Lys Thr Ala Asp Gly Leu
 225 230 235 240
 Gly Met Leu Val Gly Gln Ala Ala Ala Ser Tyr Ala Leu Trp Arg Gly
 245 250 255
 Phe Thr Pro Asp Ile Arg Pro Val Ile Glu Tyr Met Lys Ala Leu
 260 265 270

<210> 45

<211> 546

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(546)

<400> 45

ttg ctt tct cat ttg gat atg aaa ttc gtc agc gac ctt ttg tct gtc	48
Leu Leu Ser His Leu Asp Met Lys Phe Val Ser Asp Leu Leu Ser Val	
1 5 10 15	
atc ttg ttt ttt gct act tat acc gtt acc aaa aat atg att gcc gct	96
Ile Leu Phe Phe Ala Thr Tyr Thr Val Thr Lys Asn Met Ile Ala Ala	
20 25 30	
acg gcg gtt gcc ttg gtt gcc ggt gtg gtt cag gcg gct ttt ctg tat	144
Thr Ala Val Ala Leu Val Ala Gly Val Val Gln Ala Ala Phe Leu Tyr	
35 40 45	
tgg aaa tat aaa aag ctg gat acg atg cag tgg gtc gga ctg gtg ctg	192
Trp Lys Tyr Lys Lys Leu Asp Thr Met Gln Trp Val Gly Leu Val Leu	
50 55 60	
att gtc gta ttc ggc ggc gca acc att gtt ttg ggc gac agc cgc ttc	240
Ile Val Val Phe Gly Gly Ala Thr Ile Val Leu Gly Asp Ser Arg Phe	
65 70 75 80	
att atg tgg aag ccg agc gtt ttg ttt tgg ctg ggc gcg ctg ttc ctg	288
Ile Met Trp Lys Pro Ser Val Leu Phe Trp Leu Gly Ala Leu Phe Leu	
85 90 95	
tgg ggc agc cac ctc gcc ggt aaa aac ggc ttg aag gcg agt atc ggc	336
Trp Gly Ser His Leu Ala Gly Lys Asn Gly Leu Lys Ala Ser Ile Gly	
100 105 110	
agg gag att cag ctt ccg gat gcc gta tgg gcg aaa ttg acg tat atg	384
Arg Glu Ile Gln Leu Pro Asp Ala Val Trp Ala Lys Leu Thr Tyr Met	
115 120 125	
tgg gtc ggt ttc ctg att ttt atg ggt atc gcc aac tgg ttt gtg ttt	432
Trp Val Gly Phe Leu Ile Phe Met Gly Ile Ala Asn Trp Phe Val Phe	
130 135 140	
acc cgg ttc gag tcg caa tgg gtc aac tat aaa atg ttc ggc tcg act	480
Thr Arg Phe Glu Ser Gln Trp Val Asn Tyr Lys Met Phe Gly Ser Thr	
145 150 155 160	
gca ctg atg ctt gtt ttc ttt att att cag ggt att tat ctg agt acc	528
Ala Leu Met Leu Val Phe Phe Ile Ile Gln Gly Ile Tyr Leu Ser Thr	
165 170 175	
tgt ctg aaa aag gag gat	546
Cys Leu Lys Lys Glu Asp	
180	

<210> 46

<211> 182

<212> PRT

<213> Neisseria meningitidis

<400> 46

Leu Leu Ser His Leu Asp Met Lys Phe Val Ser Asp Leu Leu Ser Val
 1 5 10 15

Ile Leu Phe Phe Ala Thr Tyr Thr Val Thr Lys Asn Met Ile Ala Ala
 20 25 30

Thr Ala Val Ala Leu Val Ala Gly Val Val Gln Ala Ala Phe Leu Tyr
 35 40 45

Trp Lys Tyr Lys Lys Leu Asp Thr Met Gln Trp Val Gly Leu Val Leu
 50 55 60

Ile Val Val Phe Gly Gly Ala Thr Ile Val Leu Gly Asp Ser Arg Phe
 65 70 75 80

Ile Met Trp Lys Pro Ser Val Leu Phe Trp Leu Gly Ala Leu Phe Leu
 85 90 95

Trp Gly Ser His Leu Ala Gly Lys Asn Gly Leu Lys Ala Ser Ile Gly
 100 105 110

Arg Glu Ile Gln Leu Pro Asp Ala Val Trp Ala Lys Leu Thr Tyr Met
 115 120 125

Trp Val Gly Phe Leu Ile Phe Met Gly Ile Ala Asn Trp Phe Val Phe
 130 135 140

Thr Arg Phe Glu Ser Gln Trp Val Asn Tyr Lys Met Phe Gly Ser Thr
 145 150 155 160

Ala Leu Met Leu Val Phe Phe Ile Ile Gln Gly Ile Tyr Leu Ser Thr
 165 170 175

Cys Leu Lys Lys Glu Asp
 180

<210> 47

<211> 585

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(585)

<400> 47

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ttg aat att aaa ctg aaa acc ttg tta ttg ccc ttc gcc acg ctg gca 48
Leu Asn Ile Lys Leu Lys Thr Leu Leu Leu Pro Phe Ala Thr Leu Ala
  1             5             10             15

ttg tgc acc aat gct ttt gcc gcc ccg ccc agc gac gcg tcg ttg gcg 96
Leu Cys Thr Asn Ala Phe Ala Ala Pro Pro Ser Asp Ala Ser Leu Ala
             20             25             30

cgt tgg ctg gat acg cag aat ttt gac cgg gat ata gaa aaa aat atg 144
Arg Trp Leu Asp Thr Gln Asn Phe Asp Arg Asp Ile Glu Lys Asn Met
             35             40             45

att gag ggc ttt aat gcc gga ttt aaa ccg tat gcg gac aaa gcc ctt 192
Ile Glu Gly Phe Asn Ala Gly Phe Lys Pro Tyr Ala Asp Lys Ala Leu
             50             55             60

gcc gaa atg ccg gaa gcg aaa aaa gat cag gcg gca gaa gcc ttt aac 240
Ala Glu Met Pro Glu Ala Lys Lys Asp Gln Ala Ala Glu Ala Phe Asn
             65             70             75             80

cgt tat cgt gag aat gtt ttg aaa gat ttg att acg ccc gaa gtg aaa 288
Arg Tyr Arg Glu Asn Val Leu Lys Asp Leu Ile Thr Pro Glu Val Lys
             85             90             95

cag gct gtc cgc aat act tta ttg aag aat gcc cgt gag ata tac acg 336
Gln Ala Val Arg Asn Thr Leu Leu Lys Asn Ala Arg Glu Ile Tyr Thr
             100             105             110

caa gaa gaa att gac ggc atg att gcc ttt tac ggt tcg cct gtc ggt 384
Gln Glu Glu Ile Asp Gly Met Ile Ala Phe Tyr Gly Ser Pro Val Gly
             115             120             125

cag tcc gtc gtt gcc aaa aat ccg cgc tta atc aag aaa tcg atg agt 432
Gln Ser Val Val Ala Lys Asn Pro Arg Leu Ile Lys Lys Ser Met Ser
             130             135             140

gaa ata gcg gta tct tgg act gca ttg tca ggg aaa atc gcg caa cat 480
Glu Ile Ala Val Ser Trp Thr Ala Leu Ser Gly Lys Ile Ala Gln His
             145             150             155             160

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cat ctg ccc gag ttt acg gaa gag ttg cgg cgc atc atc tgc ggc ggt 528
 His Leu Pro Glu Phe Thr Glu Glu Leu Arg Arg Ile Ile Cys Gly Gly
 165 170 175

aaa aat ccc gat gcg ggc tgt aaa caa gcc gga cag gtt ggg aaa agg 576
 Lys Asn Pro Asp Ala Gly Cys Lys Gln Ala Gly Gln Val Gly Lys Arg
 180 185 190

cat cag aaa 585
 His Gln Lys
 195

<210> 48

<211> 195

<212> PRT

<213> Neisseria meningitidis

<400> 48

Leu Asn Ile Lys Leu Lys Thr Leu Leu Leu Pro Phe Ala Thr Leu Ala
 1 5 10 15

Leu Cys Thr Asn Ala Phe Ala Ala Pro Pro Ser Asp Ala Ser Leu Ala
 20 25 30

Arg Trp Leu Asp Thr Gln Asn Phe Asp Arg Asp Ile Glu Lys Asn Met
 35 40 45

Ile Glu Gly Phe Asn Ala Gly Phe Lys Pro Tyr Ala Asp Lys Ala Leu
 50 55 60

Ala Glu Met Pro Glu Ala Lys Lys Asp Gln Ala Ala Glu Ala Phe Asn
 65 70 75 80

Arg Tyr Arg Glu Asn Val Leu Lys Asp Leu Ile Thr Pro Glu Val Lys
 85 90 95

Gln Ala Val Arg Asn Thr Leu Leu Lys Asn Ala Arg Glu Ile Tyr Thr
 100 105 110

Gln Glu Glu Ile Asp Gly Met Ile Ala Phe Tyr Gly Ser Pro Val Gly
 115 120 125

Gln Ser Val Val Ala Lys Asn Pro Arg Leu Ile Lys Lys Ser Met Ser
 130 135 140

Glu Ile Ala Val Ser Trp Thr Ala Leu Ser Gly Lys Ile Ala Gln His
 145 150 155 160

His Leu Pro Glu Phe Thr Glu Glu Leu Arg Arg Ile Ile Cys Gly Gly
 165 170 175

Lys Asn Pro Asp Ala Gly Cys Lys Gln Ala Gly Gln Val Gly Lys Arg
 180 185 190

His Gln Lys
 195

<210> 49

<211> 462

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(462)

<400> 49

ttg ctt tgc ccg gaa aaa atg tcg ggg atg gcg gga cag tat ccg tac 48
 Leu Leu Cys Pro Glu Lys Met Ser Gly Met Ala Gly Gln Tyr Pro Tyr
 1 5 10 15

ggc gtc cgg tcg ggt ttg cgg agg aac ggc ttg aaa ctt tgg gat att 96
 Gly Val Arg Ser Gly Leu Arg Arg Asn Gly Leu Lys Leu Trp Asp Ile
 20 25 30

cat ttt aga atg acc cgt ttt atc gtc gca aga tgc ggt tta ttg ttt 144
 His Phe Arg Met Thr Arg Phe Ile Val Ala Arg Cys Gly Leu Leu Phe
 35 40 45

gca acc ctt aaa gga aaa acc atg aag aaa atg ttc gtg ctg ttc tgt 192
 Ala Thr Leu Lys Gly Lys Thr Met Lys Lys Met Phe Val Leu Phe Cys
 50 55 60

atg ctg ttc tcc tgc gcc ttc tcc ctt gcg gcg gta aac atc aat gcg 240
 Met Leu Phe Ser Cys Ala Phe Ser Leu Ala Ala Val Asn Ile Asn Ala
 65 70 75 80

gct tcg cag cag gag ctg gag gcg ctg ccg ggc ata ggc cct gcg aag 288
 Ala Ser Gln Gln Glu Leu Glu Ala Leu Pro Gly Ile Gly Pro Ala Lys
 85 90 95

gcg aag gcc att gcg gaa tac cgt gcg caa aac ggt gcg ttc aag tct 336
 Ala Lys Ala Ile Ala Glu Tyr Arg Ala Gln Asn Gly Ala Phe Lys Ser

100	105	110	
gta gac gat ttg acc aag gta aag ggc atc ggc cct gcg gtg ctg gcg			384
Val Asp Asp Leu Thr Lys Val Lys Gly Ile Gly Pro Ala Val Leu Ala			
115	120	125	
aag ctg aag gat cag gct tct gtc ggt gcg ccc gca cca aaa ggc cca			432
Lys Leu Lys Asp Gln Ala Ser Val Gly Ala Pro Ala Pro Lys Gly Pro			
130	135	140	
gct aaa cca gtg ctg ccc gcg gat aaa aaa			462
Ala Lys Pro Val Leu Pro Ala Asp Lys Lys			
145	150		
<210> 50			
<211> 154			
<212> PRT			
<213> Neisseria meningitidis			
<400> 50			
Leu Leu Cys Pro Glu Lys Met Ser Gly Met Ala Gly Gln Tyr Pro Tyr			
1	5	10	15
Gly Val Arg Ser Gly Leu Arg Arg Asn Gly Leu Lys Leu Trp Asp Ile			
20	25	30	
His Phe Arg Met Thr Arg Phe Ile Val Ala Arg Cys Gly Leu Leu Phe			
35	40	45	
Ala Thr Leu Lys Gly Lys Thr Met Lys Lys Met Phe Val Leu Phe Cys			
50	55	60	
Met Leu Phe Ser Cys Ala Phe Ser Leu Ala Ala Val Asn Ile Asn Ala			
65	70	75	80
Ala Ser Gln Gln Glu Leu Glu Ala Leu Pro Gly Ile Gly Pro Ala Lys			
85	90	95	
Ala Lys Ala Ile Ala Glu Tyr Arg Ala Gln Asn Gly Ala Phe Lys Ser			
100	105	110	
Val Asp Asp Leu Thr Lys Val Lys Gly Ile Gly Pro Ala Val Leu Ala			
115	120	125	
Lys Leu Lys Asp Gln Ala Ser Val Gly Ala Pro Ala Pro Lys Gly Pro			
130	135	140	

Ala Lys Pro Val Leu Pro Ala Asp Lys Lys
145 150

<210> 51

<211> 969

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(969)

<400> 51

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atg tcc gcc aag ttc caa caa gaa acc ctc aaa tcc cgt ttc gcg caa 48
Met Ser Ala Lys Phe Gln Gln Glu Thr Leu Lys Ser Arg Phe Ala Gln
  1             5             10             15

gcc aaa gtc ctg gtt gtc ggc gac gtg atg ctc gac cgc tat tgg ttc 96
Ala Lys Val Leu Val Val Gly Asp Val Met Leu Asp Arg Tyr Trp Phe
          20             25             30

ggc gat gtg tcc cgt att tcg ccc gaa gcc ccc gtg ccg gtg gcg aaa 144
Gly Asp Val Ser Arg Ile Ser Pro Glu Ala Pro Val Pro Val Ala Lys
          35             40             45

atc gga cga atc gac caa cgc gcg ggc gga gcg gca aat gtc gcg cgc 192
Ile Gly Arg Ile Asp Gln Arg Ala Gly Gly Ala Ala Asn Val Ala Arg
          50             55             60

aac atc gct tcg ctg ggc ggc aaa gta ggg ctg ttg tcg gta acc ggt 240
Asn Ile Ala Ser Leu Gly Gly Lys Val Gly Leu Leu Ser Val Thr Gly
          65             70             75             80

aac gac gaa gcc gcc gac gcg ctc gac gcg ctg atg gtg cag gac ggc 288
Asn Asp Glu Ala Ala Asp Ala Leu Asp Ala Leu Met Val Gln Asp Gly
          85             90             95

gtc gcc tcc tat ctg atg cgc gac aaa caa atc gcc acc acc gtc aaa 336
Val Ala Ser Tyr Leu Met Arg Asp Lys Gln Ile Ala Thr Thr Val Lys
          100            105            110

ctg cgc gtc gtc gcc cgc aac cag cag ctt atc cgc ctt gat ttt gaa 384
Leu Arg Val Val Ala Arg Asn Gln Gln Leu Ile Arg Leu Asp Phe Glu
          115            120            125

gaa cat ccc aac cgc gaa gtg ttg gaa caa atc aag cgg aaa tac cgc 432

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Glu His Pro Asn Arg Glu Val Leu Glu Gln Ile Lys Arg Lys Tyr Arg	
130 135 140	
gaa atc ttg ccc gaa tac gac gca atc att ttt tca gac tac ggc aaa	480
Glu Ile Leu Pro Glu Tyr Asp Ala Ile Ile Phe Ser Asp Tyr Gly Lys	
145 150 155 160	
ggc ggc ctg tcg cac atc tcc gat atg atc gat tgg gcg aaa cac gaa	528
Gly Gly Leu Ser His Ile Ser Asp Met Ile Asp Trp Ala Lys His Glu	
165 170 175	
ggc aaa acc gta tta atc gac ccc aaa ggc gac gat tac gaa aaa tac	576
Gly Lys Thr Val Leu Ile Asp Pro Lys Gly Asp Asp Tyr Glu Lys Tyr	
180 185 190	
gcc ggc gca acc ctg att acg ccc aac cgc gcc gaa ttg aaa gaa gtg	624
Ala Gly Ala Thr Leu Ile Thr Pro Asn Arg Ala Glu Leu Lys Glu Val	
195 200 205	
gtc ggc agc tgg aaa aac gaa aat gat ttg acc gaa aaa gcc caa aac	672
Val Gly Ser Trp Lys Asn Glu Asn Asp Leu Thr Glu Lys Ala Gln Asn	
210 215 220	
ctg cgc cgc cac ctc gac ttg acc gct att tta ctg acc cga agc gaa	720
Leu Arg Arg His Leu Asp Leu Thr Ala Ile Leu Leu Thr Arg Ser Glu	
225 230 235 240	
gaa ggc atg acc ttg ttc agc gaa ggc gaa ccc att tac cag ccc acc	768
Glu Gly Met Thr Leu Phe Ser Glu Gly Glu Pro Ile Tyr Gln Pro Thr	
245 250 255	
cgc gcc caa gaa gtt tac gac gta tcc ggt gcg ggc gac acc gtc att	816
Arg Ala Gln Glu Val Tyr Asp Val Ser Gly Ala Gly Asp Thr Val Ile	
260 265 270	
gcc gga atg ggc ttg ggt ttg gcg gca ggc tgc acc atg ccc gaa gcc	864
Ala Gly Met Gly Leu Gly Leu Ala Ala Gly Cys Thr Met Pro Glu Ala	
275 280 285	
atg tac ctt gcc aat act gcg gcc ggg gtt gtc gtg gcg aaa ctc ggt	912
Met Tyr Leu Ala Asn Thr Ala Ala Gly Val Val Val Ala Lys Leu Gly	
290 295 300	
acg gcg gtt tgc tcg ttt gcc gaa ttg acc aag gca ttg tca ggg caa	960
Thr Ala Val Cys Ser Phe Ala Glu Leu Thr Lys Ala Leu Ser Gly Gln	
305 310 315 320	
tca aca atg	969

Ser Thr Met

<210> 52

<211> 323

<212> PRT

<213> Neisseria meningitidis

<400> 52

Met Ser Ala Lys Phe Gln Gln Glu Thr Leu Lys Ser Arg Phe Ala Gln

1 5 10 15

Ala Lys Val Leu Val Val Gly Asp Val Met Leu Asp Arg Tyr Trp Phe

20 25 30

Gly Asp Val Ser Arg Ile Ser Pro Glu Ala Pro Val Pro Val Ala Lys

35 40 45

Ile Gly Arg Ile Asp Gln Arg Ala Gly Gly Ala Ala Asn Val Ala Arg

50 55 60

Asn Ile Ala Ser Leu Gly Gly Lys Val Gly Leu Leu Ser Val Thr Gly

65 70 75 80

Asn Asp Glu Ala Ala Asp Ala Leu Asp Ala Leu Met Val Gln Asp Gly

85 90 95

Val Ala Ser Tyr Leu Met Arg Asp Lys Gln Ile Ala Thr Thr Val Lys

100 105 110

Leu Arg Val Val Ala Arg Asn Gln Gln Leu Ile Arg Leu Asp Phe Glu

115 120 125

Glu His Pro Asn Arg Glu Val Leu Glu Gln Ile Lys Arg Lys Tyr Arg

130 135 140

Glu Ile Leu Pro Glu Tyr Asp Ala Ile Ile Phe Ser Asp Tyr Gly Lys

145 150 155 160

Gly Gly Leu Ser His Ile Ser Asp Met Ile Asp Trp Ala Lys His Glu

165 170 175

Gly Lys Thr Val Leu Ile Asp Pro Lys Gly Asp Asp Tyr Glu Lys Tyr

180 185 190

Ala Gly Ala Thr Leu Ile Thr Pro Asn Arg Ala Glu Leu Lys Glu Val

195 200 205

Val Gly Ser Trp Lys Asn Glu Asn Asp Leu Thr Glu Lys Ala Gln Asn
 210 215 220

Leu Arg Arg His Leu Asp Leu Thr Ala Ile Leu Leu Thr Arg Ser Glu
 225 230 235 240

Glu Gly Met Thr Leu Phe Ser Glu Gly Glu Pro Ile Tyr Gln Pro Thr
 245 250 255

Arg Ala Gln Glu Val Tyr Asp Val Ser Gly Ala Gly Asp Thr Val Ile
 260 265 270

Ala Gly Met Gly Leu Gly Leu Ala Ala Gly Cys Thr Met Pro Glu Ala
 275 280 285

Met Tyr Leu Ala Asn Thr Ala Ala Gly Val Val Val Ala Lys Leu Gly
 290 295 300

Thr Ala Val Cys Ser Phe Ala Glu Leu Thr Lys Ala Leu Ser Gly Gln
 305 310 315 320

Ser Thr Met

<210> 53
 <211> 864
 <212> DNA
 <213> Neisseria meningitidis

<220>
 <221> CDS
 <222> (1)..(864)

<400> 53
 atg aaa gca aaa atc ctg act tcc gtt gca ctg ctt gcc tgt tcc ggc 48
 Met Lys Ala Lys Ile Leu Thr Ser Val Ala Leu Leu Ala Cys Ser Gly
 1 5 10 15

agc ctg ttt gcc caa acg ctg gca acc gtc aac ggt cag aaa atc gac 96
 Ser Leu Phe Ala Gln Thr Leu Ala Thr Val Asn Gly Gln Lys Ile Asp
 20 25 30

agt tcc gtc att gat gcg cag gtt gcc gca ttc cgt gcg gaa aac agc 144
 Ser Ser Val Ile Asp Ala Gln Val Ala Ala Phe Arg Ala Glu Asn Ser
 35 40 45

cgt gcc gaa gac acg ccg caa ctg cgc caa tcc ctg ctg gaa aac gaa	192
Arg Ala Glu Asp Thr Pro Gln Leu Arg Gln Ser Leu Leu Glu Asn Glu	
50 55 60	
gtg gtc aac acc gtg gtc gca cag gaa gtg aaa cgc ctg aaa ctc gac	240
Val Val Asn Thr Val Val Ala Gln Glu Val Lys Arg Leu Lys Leu Asp	
65 70 75 80	
cgg tcg gca gag ttt aaa aat gcg ctt gcc aaa ttg cgt gcc gaa gcg	288
Arg Ser Ala Glu Phe Lys Asn Ala Leu Ala Lys Leu Arg Ala Glu Ala	
85 90 95	
aaa aag tcg ggc gac gac aag aaa ccg tcc ttc aaa acc gtt tgg cag	336
Lys Lys Ser Gly Asp Asp Lys Lys Pro Ser Phe Lys Thr Val Trp Gln	
100 105 110	
gcg gta aaa tat ggc ttg aac ggc gag gca tac gcg ctg cat atc gcc	384
Ala Val Lys Tyr Gly Leu Asn Gly Glu Ala Tyr Ala Leu His Ile Ala	
115 120 125	
aaa acc caa ccg gtt tcc gag cag gaa gta aaa gcc gca tat gac aat	432
Lys Thr Gln Pro Val Ser Glu Gln Glu Val Lys Ala Ala Tyr Asp Asn	
130 135 140	
atc agc ggt ttt tac aaa ggt acg cag gaa gtc cag ttg ggc gaa atc	480
Ile Ser Gly Phe Tyr Lys Gly Thr Gln Glu Val Gln Leu Gly Glu Ile	
145 150 155 160	
ctg acc gac aag gaa gaa aat gca aaa aaa gcg gtt gcc gac ttg aag	528
Leu Thr Asp Lys Glu Glu Asn Ala Lys Lys Ala Val Ala Asp Leu Lys	
165 170 175	
gcg aaa aaa ggt ttc gat gcc gtc ttg aaa caa tat tcc ctc aac gac	576
Ala Lys Lys Gly Phe Asp Ala Val Leu Lys Gln Tyr Ser Leu Asn Asp	
180 185 190	
cgt acc aaa cag acc ggt gcg ccg gtc gga tat gtg ccg ctg aaa gat	624
Arg Thr Lys Gln Thr Gly Ala Pro Val Gly Tyr Val Pro Leu Lys Asp	
195 200 205	
ttg gaa cag ggt gtt ccg ccg ctt tat cag gca att aag gac ttg aaa	672
Leu Glu Gln Gly Val Pro Pro Leu Tyr Gln Ala Ile Lys Asp Leu Lys	
210 215 220	
aaa ggc gaa ttt acg gca acg ccg ctg aaa aac ggc gat ttc tac ggc	720
Lys Gly Glu Phe Thr Ala Thr Pro Leu Lys Asn Gly Asp Phe Tyr Gly	
225 230 235 240	

gtt tat tat gtc aac gac agc cgc gag gta aaa gtg cct tct ttt gat 768
 Val Tyr Tyr Val Asn Asp Ser Arg Glu Val Lys Val Pro Ser Phe Asp
 245 250 255

gaa atg aaa gga cag att gcg ggc aac ctt cag gcg gaa cgg att gac 816
 Glu Met Lys Gly Gln Ile Ala Gly Asn Leu Gln Ala Glu Arg Ile Asp
 260 265 270

cgt gcc gtc ggt gca ctg ttg ggc aag gca aac atc aaa cct gca aaa 864
 Arg Ala Val Gly Ala Leu Leu Gly Lys Ala Asn Ile Lys Pro Ala Lys
 275 280 285

<210> 54

<211> 288

<212> PRT

<213> Neisseria meningitidis

<400> 54

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Ser Leu Phe Ala Gln Thr Leu Ala Thr Val Asn Gly Gln Lys Ile Asp
 20 25 30

Ser Ser Val Ile Asp Ala Gln Val Ala Ala Phe Arg Ala Glu Asn Ser
 35 40 45

Arg Ala Glu Asp Thr Pro Gln Leu Arg Gln Ser Leu Leu Glu Asn Glu
 50 55 60

Val Val Asn Thr Val Val Ala Gln Glu Val Lys Arg Leu Lys Leu Asp
 65 70 75 80

Arg Ser Ala Glu Phe Lys Asn Ala Leu Ala Lys Leu Arg Ala Glu Ala
 85 90 95

Lys Lys Ser Gly Asp Asp Lys Lys Pro Ser Phe Lys Thr Val Trp Gln
 100 105 110

Ala Val Lys Tyr Gly Leu Asn Gly Glu Ala Tyr Ala Leu His Ile Ala
 115 120 125

Lys Thr Gln Pro Val Ser Glu Gln Glu Val Lys Ala Ala Tyr Asp Asn
 130 135 140

Ile Ser Gly Phe Tyr Lys Gly Thr Gln Glu Val Gln Leu Gly Glu Ile
 145 150 155 160

Leu Thr Asp Lys Glu Glu Asn Ala Lys Lys Ala Val Ala Asp Leu Lys
 165 170 175

Ala Lys Lys Gly Phe Asp Ala Val Leu Lys Gln Tyr Ser Leu Asn Asp
 180 185 190

Arg Thr Lys Gln Thr Gly Ala Pro Val Gly Tyr Val Pro Leu Lys Asp
 195 200 205

Leu Glu Gln Gly Val Pro Pro Leu Tyr Gln Ala Ile Lys Asp Leu Lys
 210 215 220

Lys Gly Glu Phe Thr Ala Thr Pro Leu Lys Asn Gly Asp Phe Tyr Gly
 225 230 235 240

Val Tyr Tyr Val Asn Asp Ser Arg Glu Val Lys Val Pro Ser Phe Asp
 245 250 255

Glu Met Lys Gly Gln Ile Ala Gly Asn Leu Gln Ala Glu Arg Ile Asp
 260 265 270

Arg Ala Val Gly Ala Leu Leu Gly Lys Ala Asn Ile Lys Pro Ala Lys
 275 280 285

<210> 55

<211> 1257

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1257)

<400> 55

atg aaa cag acc gtc ctc aaa aat aac ctg caa aac ctg ctt gaa agc 48
 Met Lys Gln Thr Val Leu Lys Asn Asn Leu Gln Asn Leu Leu Glu Ser
 1 5 10 15

gca gaa aat atc ctg ctg ctt caa ggc cct gtc ggc gat ttt ttt ctg 96
 Ala Glu Asn Ile Leu Leu Leu Gln Gly Pro Val Gly Asp Phe Phe Leu
 20 25 30

cgc ctt gcc gac tgg ctg act gca aac ggc aaa acc gta cat aaa ttc 144
 Arg Leu Ala Asp Trp Leu Thr Ala Asn Gly Lys Thr Val His Lys Phe
 35 40 45


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aac ttt aat gca ggc gac gac tat ttt tat ccg ccc act caa gcg cat 192
Asn Phe Asn Ala Gly Asp Asp Tyr Phe Tyr Pro Pro Thr Gln Ala His
    50                55                60

acc gtt gtt ttt aac gac aac tac gat gcc ttt cct gag ttt ttg caa 240
Thr Val Val Phe Asn Asp Asn Tyr Asp Ala Phe Pro Glu Phe Leu Gln
    65                70                75                80

gaa tac atc act caa cat cac atc cag gcc gtt gtc tgc ttt ggc gac 288
Glu Tyr Ile Thr Gln His His Ile Gln Ala Val Val Cys Phe Gly Asp
                85                90                95

aca cgc cct tat cac gtc att gca aaa cgc att gca aac gaa aac caa 336
Thr Arg Pro Tyr His Val Ile Ala Lys Arg Ile Ala Asn Glu Asn Gln
                100                105                110

gcc agt ttc tgg gcg ttt gaa gaa ggc tat ttc cgc ccc tac tac atc 384
Ala Ser Phe Trp Ala Phe Glu Glu Gly Tyr Phe Arg Pro Tyr Tyr Ile
                115                120                125

acc tta gaa aaa gac ggc gtc aac gca ttt tcc ccg ttg ccg cgc cgt 432
Thr Leu Glu Lys Asp Gly Val Asn Ala Phe Ser Pro Leu Pro Arg Arg
                130                135                140

gcc gac ttt ttt ctt gaa caa ttc cct aag ctt gcc cag caa gaa tat 480
Ala Asp Phe Phe Leu Glu Gln Phe Pro Lys Leu Ala Gln Gln Glu Tyr
                145                150                155                160

aaa gcg cca acg ccg gta cac ggc ggt ttt acg ccc atg gca aaa aac 528
Lys Ala Pro Thr Pro Val His Gly Gly Phe Thr Pro Met Ala Lys Asn
                165                170                175

gct atc cgt tac tat atc gag ttg ttc cgc aat cta cgc aaa tac ccc 576
Ala Ile Arg Tyr Tyr Ile Glu Leu Phe Arg Asn Leu Arg Lys Tyr Pro
                180                185                190

gac tac atc cac cac cgc gca ccc aat gcc ggc cat tac ctc aaa ccg 624
Asp Tyr Ile His His Arg Ala Pro Asn Ala Gly His Tyr Leu Lys Pro
                195                200                205

tgg tcg ctc tcc atc ctc aag cgt ttg aac tac tat att gaa gac atc 672
Trp Ser Leu Ser Ile Leu Lys Arg Leu Asn Tyr Tyr Ile Glu Asp Ile
                210                215                220

caa atc gcc aaa cgt gtg gaa gca ggc aaa tac ggc aag ttt ttt att 720
Gln Ile Ala Lys Arg Val Glu Ala Gly Lys Tyr Gly Lys Phe Phe Ile
                225                230                235                240

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gtt ccc ttg cag gta ttc aac gac agc caa gtc cgt atc cat tgc gac	768
Val Pro Leu Gln Val Phe Asn Asp Ser Gln Val Arg Ile His Cys Asp	
245 250 255	
ttt ccc agc gtc cgc agc ttc ctg ctc cat gtt ttg agt tca ttt gcc	816
Phe Pro Ser Val Arg Ser Phe Leu Leu His Val Leu Ser Ser Phe Ala	
260 265 270	
gaa cac gcg cct gcc gat acc aac atc atc atc aag cat cat cct atg	864
Glu His Ala Pro Ala Asp Thr Asn Ile Ile Ile Lys His His Pro Met	
275 280 285	
gac cgc ggc ttt atc gac tac tgg cgc gac att aaa cgc ttt atc aaa	912
Asp Arg Gly Phe Ile Asp Tyr Trp Arg Asp Ile Lys Arg Phe Ile Lys	
290 295 300	
gaa cac ccc gaa ctc aaa ggc cgt gtg att tat gtc cat gat gtc ccc	960
Glu His Pro Glu Leu Lys Gly Arg Val Ile Tyr Val His Asp Val Pro	
305 310 315 320	
ctg ccc gtt ttc ctg cgc cac ggt ctc ggc atg gtt acc atc aac agc	1008
Leu Pro Val Phe Leu Arg His Gly Leu Gly Met Val Thr Ile Asn Ser	
325 330 335	
acc agc ggc ctg tcc gga ctg att cac aat atg cca gtt aag gtt ctc	1056
Thr Ser Gly Leu Ser Gly Leu Ile His Asn Met Pro Val Lys Val Leu	
340 345 350	
ggc cgt gcc tat tat gat att ccc ggc att act gac caa aat acc ttg	1104
Gly Arg Ala Tyr Tyr Asp Ile Pro Gly Ile Thr Asp Gln Asn Thr Leu	
355 360 365	
gca gaa ttt tgg aat cat ccg aca ccg cct gac aaa gag ctg ttc cat	1152
Ala Glu Phe Trp Asn His Pro Thr Pro Pro Asp Lys Glu Leu Phe His	
370 375 380	
gcc tac cga atg tac cac ctc aac gtg acc caa att aac ggc aac ttc	1200
Ala Tyr Arg Met Tyr His Leu Asn Val Thr Gln Ile Asn Gly Asn Phe	
385 390 395 400	
tac agt cag gtc ttt ttt cct aac aaa aac aca tcc gac tct tcc aca	1248
Tyr Ser Gln Val Phe Phe Pro Asn Lys Asn Thr Ser Asp Ser Ser Thr	
405 410 415	
cca aca acc	1257
Pro Thr Thr	

<210> 56

<211> 419

<212> PRT

<213> Neisseria meningitidis

<400> 56

Met Lys Gln Thr Val Leu Lys Asn Asn Leu Gln Asn Leu Leu Glu Ser
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Ala Glu Asn Ile Leu Leu Leu Gln Gly Pro Val Gly Asp Phe Phe Leu
 20 25 30

Arg Leu Ala Asp Trp Leu Thr Ala Asn Gly Lys Thr Val His Lys Phe
 35 40 45

Asn Phe Asn Ala Gly Asp Asp Tyr Phe Tyr Pro Pro Thr Gln Ala His
 50 55 60

Thr Val Val Phe Asn Asp Asn Tyr Asp Ala Phe Pro Glu Phe Leu Gln
 65 70 75 80

Glu Tyr Ile Thr Gln His His Ile Gln Ala Val Val Cys Phe Gly Asp
 85 90 95

Thr Arg Pro Tyr His Val Ile Ala Lys Arg Ile Ala Asn Glu Asn Gln
 100 105 110

Ala Ser Phe Trp Ala Phe Glu Glu Gly Tyr Phe Arg Pro Tyr Tyr Ile
 115 120 125

Thr Leu Glu Lys Asp Gly Val Asn Ala Phe Ser Pro Leu Pro Arg Arg
 130 135 140

Ala Asp Phe Phe Leu Glu Gln Phe Pro Lys Leu Ala Gln Gln Glu Tyr
 145 150 155 160

Lys Ala Pro Thr Pro Val His Gly Gly Phe Thr Pro Met Ala Lys Asn
 165 170 175

Ala Ile Arg Tyr Tyr Ile Glu Leu Phe Arg Asn Leu Arg Lys Tyr Pro
 180 185 190

Asp Tyr Ile His His Arg Ala Pro Asn Ala Gly His Tyr Leu Lys Pro
 195 200 205

Trp Ser Leu Ser Ile Leu Lys Arg Leu Asn Tyr Tyr Ile Glu Asp Ile
 210 215 220

Gln Ile Ala Lys Arg Val Glu Ala Gly Lys Tyr Gly Lys Phe Phe Ile
 225 230 235 240

Val Pro Leu Gln Val Phe Asn Asp Ser Gln Val Arg Ile His Cys Asp
 245 250 255

Phe Pro Ser Val Arg Ser Phe Leu Leu His Val Leu Ser Ser Phe Ala
 260 265 270

Glu His Ala Pro Ala Asp Thr Asn Ile Ile Ile Lys His His Pro Met
 275 280 285

Asp Arg Gly Phe Ile Asp Tyr Trp Arg Asp Ile Lys Arg Phe Ile Lys
 290 295 300

Glu His Pro Glu Leu Lys Gly Arg Val Ile Tyr Val His Asp Val Pro
 305 310 315 320

Leu Pro Val Phe Leu Arg His Gly Leu Gly Met Val Thr Ile Asn Ser
 325 330 335

Thr Ser Gly Leu Ser Gly Leu Ile His Asn Met Pro Val Lys Val Leu
 340 345 350

Gly Arg Ala Tyr Tyr Asp Ile Pro Gly Ile Thr Asp Gln Asn Thr Leu
 355 360 365

Ala Glu Phe Trp Asn His Pro Thr Pro Pro Asp Lys Glu Leu Phe His
 370 375 380

Ala Tyr Arg Met Tyr His Leu Asn Val Thr Gln Ile Asn Gly Asn Phe
 385 390 395 400

Tyr Ser Gln Val Phe Phe Pro Asn Lys Asn Thr Ser Asp Ser Ser Thr
 405 410 415

Pro Thr Thr

<210> 57

<211> 1407

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1) .. (1407)

<400> 57

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Met Thr Ser Thr Phe Pro Arg Arg Leu Ala Arg Lys Ile Arg Gln Thr
  1             5             10            15

cgc cgc ctg tgc cgc aaa agc atc gcc ttt ctg ttc ctt ttg gca ggt 96
Arg Arg Leu Ser Arg Lys Ser Ile Ala Phe Leu Phe Leu Leu Ala Gly
          20             25             30

tcg gca ctc gtc gcc ctg acc gcg ctg ttt ttt gcc cat ctt gcc gat 144
Ser Ala Leu Val Ala Leu Thr Ala Leu Phe Phe Ala His Leu Ala Asp
        35             40             45

ttt gcg ctg gaa ctg aac gcc aaa ctg gtt caa caa tac ccg tgg ttc 192
Phe Ala Leu Glu Leu Asn Ala Lys Leu Val Gln Gln Tyr Pro Trp Phe
        50             55             60

gcg tgg gtc gcg ctt cct ttg ggt tta ccg ctt att gcg tgg ctc aca 240
Ala Trp Val Ala Leu Pro Leu Gly Leu Pro Leu Ile Ala Trp Leu Thr
        65             70             75             80

cgc aaa ttc gcc ccc ttc acc gcc ggc agc ggc atc ccg cag gtc atc 288
Arg Lys Phe Ala Pro Phe Thr Ala Gly Ser Gly Ile Pro Gln Val Ile
          85             90             95

gcc tca ctg tgc ctg ccc tac ggc gca cag aaa acg cgg ctg atc cgc 336
Ala Ser Leu Ser Leu Pro Tyr Gly Ala Gln Lys Thr Arg Leu Ile Arg
        100            105            110

ctc ggg cag acg ctg ctg aag att ccg cta acc ttt ttg ggt atg ctg 384
Leu Gly Gln Thr Leu Leu Lys Ile Pro Leu Thr Phe Leu Gly Met Leu
        115            120            125

ttc ggc gcg tcc atc gga cgc gaa ggt ccg tcc gtg cag gtc ggc gcg 432
Phe Gly Ala Ser Ile Gly Arg Glu Gly Pro Ser Val Gln Val Gly Ala
        130            135            140

gca gtg atg ggc gcg tgg ggc gcg tgg tgc aag aaa cac ggc ttg gca 480
Ala Val Met Gly Ala Trp Gly Ala Trp Cys Lys Lys His Gly Leu Ala
        145            150            155            160

ttc aaa ggg atg cag gaa aac gat ttg atg gcg gcg ggc gcg gcg ggc 528
Phe Lys Gly Met Gln Glu Asn Asp Leu Met Ala Ala Gly Ala Ala Gly
        165            170            175

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ggt ttg gca gcc gcg ttc aac gcg ccg ctg gcg ggc gtg att ttc gcc	576
Gly Leu Ala Ala Ala Phe Asn Ala Pro Leu Ala Gly Val Ile Phe Ala	
180 185 190	
att gag gaa ctc ggg cgc ggc atc atg ttg cgc tgg gag agg caa att	624
Ile Glu Glu Leu Gly Arg Gly Ile Met Leu Arg Trp Glu Arg Gln Ile	
195 200 205	
ctt ttg ggc gtg ctc gcc tcc ggt ttc ata cag gtc gcc att cag ggc	672
Leu Leu Gly Val Leu Ala Ser Gly Phe Ile Gln Val Ala Ile Gln Gly	
210 215 220	
aac aac ccg tat ttt tcc ggc ttc aac ggc ggc gta ttg gaa cat atc	720
Asn Asn Pro Tyr Phe Ser Gly Phe Asn Gly Gly Val Leu Glu His Ile	
225 230 235 240	
ttt ctg tgg gtc gca ctg tcc ggc ctg gtt tgc ggc gcg gcg ggc ggg	768
Phe Leu Trp Val Ala Leu Ser Gly Leu Val Cys Gly Ala Ala Gly Gly	
245 250 255	
ctg ttc gga cgt ttg ctc tat cgc ggt gcg gcg gcg ttt gca ccg cgc	816
Leu Phe Gly Arg Leu Leu Tyr Arg Gly Ala Ala Ala Phe Ala Pro Arg	
260 265 270	
aag ata cgc ggc ttc atc cgc aac cgt ccg ctg ctg ctg gcg gca ctg	864
Lys Ile Arg Gly Phe Ile Arg Asn Arg Pro Leu Leu Leu Ala Ala Leu	
275 280 285	
atg ggg ctg ctg ctc gcc ctg ctc ggc acg ttc tac caa ggc aaa acc	912
Met Gly Leu Leu Leu Ala Leu Leu Gly Thr Phe Tyr Gln Gly Lys Thr	
290 295 300	
tac ggc acc ggc tac cac gaa gcc gcc caa gcc ctg cac ggc atc tac	960
Tyr Gly Thr Gly Tyr His Glu Ala Ala Gln Ala Leu His Gly Ile Tyr	
305 310 315 320	
gaa gcc ccc ttc gga ctc gcc gcc gcc aaa tgg ctc gcc acc gta ttc	1008
Glu Ala Pro Phe Gly Leu Ala Ala Ala Lys Trp Leu Ala Thr Val Phe	
325 330 335	
agc tat tgg gca ggc gtt ccg ggc ggc att ttc act ccc tcg ctg acc	1056
Ser Tyr Trp Ala Gly Val Pro Gly Gly Ile Phe Thr Pro Ser Leu Thr	
340 345 350	
ata ggc gcg gtt ttg ggc gag cat atc gcc gcc atc gcc gac ata tcg	1104
Ile Gly Ala Val Leu Gly Glu His Ile Ala Ala Ile Ala Asp Ile Ser	
355 360 365	

cag ggt gca aac atc atc gtc ctc atc tgc atg gcg gca ttt ctg gcg 1152
 Gln Gly Ala Asn Ile Ile Val Leu Ile Cys Met Ala Ala Phe Leu Ala
 370 375 380

ggc gcg aca caa tcc ccg att act tcc gcc gtc gtc gtc atg gaa atg 1200
 Gly Ala Thr Gln Ser Pro Ile Thr Ser Ala Val Val Val Met Glu Met
 385 390 395 400

acg ggc gga caa agc ctg ctg ttt tgg atg ctg att gcc tgc att ttc 1248
 Thr Gly Gly Gln Ser Leu Leu Phe Trp Met Leu Ile Ala Cys Ile Phe
 405 410 415

gcc tcg cag gtt tcg cgc cag ttt tcg ccg cgt ccg ttc tac cac gca 1296
 Ala Ser Gln Val Ser Arg Gln Phe Ser Pro Arg Pro Phe Tyr His Ala
 420 425 430

tcg gga atg cgc ttc cgc cag cgc gtg ctt caa gaa acc gcc gcc caa 1344
 Ser Gly Met Arg Phe Arg Gln Arg Val Leu Gln Glu Thr Ala Ala Gln
 435 440 445

acc ggc aat gcg ccc gca aga ccg caa aca gca aac agc aaa acg gga 1392
 Thr Gly Asn Ala Pro Ala Arg Pro Gln Thr Ala Asn Ser Lys Thr Gly
 450 455 460

atg ccg tct gaa aat 1407
 Met Pro Ser Glu Asn
 465

<210> 58

<211> 469

<212> PRT

<213> Neisseria meningitidis

<400> 58

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Arg Arg Leu Ser Arg Lys Ser Ile Ala Phe Leu Phe Leu Leu Ala Gly
 20 25 30

Ser Ala Leu Val Ala Leu Thr Ala Leu Phe Phe Ala His Leu Ala Asp
 35 40 45

Phe Ala Leu Glu Leu Asn Ala Lys Leu Val Gln Gln Tyr Pro Trp Phe
 50 55 60

Ala Trp Val Ala Leu Pro Leu Gly Leu Pro Leu Ile Ala Trp Leu Thr

65	70	75	80
Arg Lys Phe Ala Pro Phe Thr Ala Gly Ser Gly Ile Pro Gln Val Ile			
	85	90	95
Ala Ser Leu Ser Leu Pro Tyr Gly Ala Gln Lys Thr Arg Leu Ile Arg			
	100	105	110
Leu Gly Gln Thr Leu Leu Lys Ile Pro Leu Thr Phe Leu Gly Met Leu			
	115	120	125
Phe Gly Ala Ser Ile Gly Arg Glu Gly Pro Ser Val Gln Val Gly Ala			
	130	135	140
Ala Val Met Gly Ala Trp Gly Ala Trp Cys Lys Lys His Gly Leu Ala			
	145	150	155
Phe Lys Gly Met Gln Glu Asn Asp Leu Met Ala Ala Gly Ala Ala Gly			
	165	170	175
Gly Leu Ala Ala Ala Phe Asn Ala Pro Leu Ala Gly Val Ile Phe Ala			
	180	185	190
Ile Glu Glu Leu Gly Arg Gly Ile Met Leu Arg Trp Glu Arg Gln Ile			
	195	200	205
Leu Leu Gly Val Leu Ala Ser Gly Phe Ile Gln Val Ala Ile Gln Gly			
	210	215	220
Asn Asn Pro Tyr Phe Ser Gly Phe Asn Gly Gly Val Leu Glu His Ile			
	225	230	235
Phe Leu Trp Val Ala Leu Ser Gly Leu Val Cys Gly Ala Ala Gly Gly			
	245	250	255
Leu Phe Gly Arg Leu Leu Tyr Arg Gly Ala Ala Ala Phe Ala Pro Arg			
	260	265	270
Lys Ile Arg Gly Phe Ile Arg Asn Arg Pro Leu Leu Leu Ala Ala Leu			
	275	280	285
Met Gly Leu Leu Leu Ala Leu Leu Gly Thr Phe Tyr Gln Gly Lys Thr			
	290	295	300
Tyr Gly Thr Gly Tyr His Glu Ala Ala Gln Ala Leu His Gly Ile Tyr			
	305	310	315
Glu Ala Pro Phe Gly Leu Ala Ala Ala Lys Trp Leu Ala Thr Val Phe			

<400> 59

atg aca caa gaa acc gct ttg ggc gca gca ctg aaa tcc gcc gtc caa 48
Met Thr Gln Glu Thr Ala Leu Gly Ala Ala Leu Lys Ser Ala Val Gln
1 5 10 15

act atg agc aaa aag aaa cag aca gaa atg att gcc gac cac atc tac 96
Thr Met Ser Lys Lys Lys Gln Thr Glu Met Ile Ala Asp His Ile Tyr
20 25 30

ggc aaa tat gat gta ttc aaa cga ttt aaa ccg ttg gca ctc ggt atc 144
 Gly Lys Tyr Asp Val Phe Lys Arg Phe Lys Pro Leu Ala Leu Gly Ile
 35 40 45

gat cag gat ttg att gcc gct ttg ccc caa tac gat gcc gca ctg att 192
 Asp Gln Asp Leu Ile Ala Ala Leu Pro Gln Tyr Asp Ala Ala Leu Ile
 50 55 60

gca cgc gtc ctc gcc aac cac tgc cgc cgt ccg cgc tat ctg aaa gcc 240
 Ala Arg Val Leu Ala Asn His Cys Arg Arg Pro Arg Tyr Leu Lys Ala
 65 70 75 80

ttg gcg cgc ggt ggc aaa cgt ttc gat ttg aac aac cgt ttc aaa ggc 288
 Leu Ala Arg Gly Gly Lys Arg Phe Asp Leu Asn Asn Arg Phe Lys Gly
 85 90 95

gaa gtt acc ccc gaa gaa cag gca atc gcg caa aac cat cct ttc gtg 336
 Glu Val Thr Pro Glu Glu Gln Ala Ile Ala Gln Asn His Pro Phe Val
 100 105 110

cag cag gct ttg caa cag cag tcc gcc caa gct gtc gcc gaa acg ccg 384
 Gln Gln Ala Leu Gln Gln Gln Ser Ala Gln Ala Val Ala Glu Thr Pro
 115 120 125

tct gtt gaa gcc gaa gca gcc gaa tct tcc aca aca gaa 423
 Ser Val Glu Ala Glu Ala Ala Glu Ser Ser Thr Thr Glu
 130 135 140

<210> 60

<211> 141

<212> PRT

<213> Neisseria meningitidis

<400> 60

Met Thr Gln Glu Thr Ala Leu Gly Ala Ala Leu Lys Ser Ala Val Gln
 1 5 10 15

Thr Met Ser Lys Lys Lys Gln Thr Glu Met Ile Ala Asp His Ile Tyr
 20 25 30

Gly Lys Tyr Asp Val Phe Lys Arg Phe Lys Pro Leu Ala Leu Gly Ile
 35 40 45

Asp Gln Asp Leu Ile Ala Ala Leu Pro Gln Tyr Asp Ala Ala Leu Ile
 50 55 60

Ala Arg Val Leu Ala Asn His Cys Arg Arg Pro Arg Tyr Leu Lys Ala
65 70 75 80

Leu Ala Arg Gly Gly Lys Arg Phe Asp Leu Asn Asn Arg Phe Lys Gly
85 90 95

Glu Val Thr Pro Glu Glu Gln Ala Ile Ala Gln Asn His Pro Phe Val
100 105 110

Gln Gln Ala Leu Gln Gln Gln Ser Ala Gln Ala Val Ala Glu Thr Pro
115 120 125

Ser Val Glu Ala Glu Ala Ala Glu Ser Ser Thr Thr Glu
130 135 140

<210> 61

<211> 1377

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1377)

<400> 61

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1 5 10 15

gcc gtc gtc gtc gct ctg att gcc gcg atg ggc tat acc atc att tca 96
Ala Val Val Val Ala Leu Ile Ala Ala Met Gly Tyr Thr Ile Ile Ser
20 25 30

ttg gag tgg ctg ccg cat atg tcc att att gcc gcc atc gtc gtg ctg 144
Leu Glu Trp Leu Pro His Met Ser Ile Ile Ala Ala Ile Val Val Leu
35 40 45

att ttg tac ggc ttg gcg cgc ggt ttg aaa tac aac gat atg cag cag 192
Ile Leu Tyr Gly Leu Ala Arg Gly Leu Lys Tyr Asn Asp Met Gln Gln
50 55 60

ggc atg ata ggc gcg ttg aat cag ggt atg ggc gcg att tac ctg ttt 240
Gly Met Ile Gly Ala Leu Asn Gln Gly Met Gly Ala Ile Tyr Leu Phe
65 70 75 80

ttc ttc atc ggg ctg atg gtc agc gcg ctg atg atg agc ggc gcg att 288

Phe Phe Ile Gly Leu Met Val Ser Ala Leu Met Met Ser Gly Ala Ile	
85 90 95	
ccg acg ctg atg tat tac ggt ttc gga ctg att tcc ccg act tat ttt	336
Pro Thr Leu Met Tyr Tyr Gly Phe Gly Leu Ile Ser Pro Thr Tyr Phe	
100 105 110	
tat ttt tcc gcc ttc gcg ctg tgt tcc gtc atc ggc gtg tcc atc ggc	384
Tyr Phe Ser Ala Phe Ala Leu Cys Ser Val Ile Gly Val Ser Ile Gly	
115 120 125	
agc agc ctg acc acc tgc gcc act gtc ggc gtt gcc ttt atg ggt atg	432
Ser Ser Leu Thr Thr Cys Ala Thr Val Gly Val Ala Phe Met Gly Met	
130 135 140	
gcg gcg gcg ttt cag gcc gat atg gcg atg acg gcg ggc gcg att gtt	480
Ala Ala Ala Phe Gln Ala Asp Met Ala Met Thr Ala Gly Ala Ile Val	
145 150 155 160	
tcg ggc gca ttt ttt ggc gac aaa atg tcc ccg ctt tcc gat acg acg	528
Ser Gly Ala Phe Phe Gly Asp Lys Met Ser Pro Leu Ser Asp Thr Thr	
165 170 175	
ggc att tcc gcg tcc att gtc ggt atc gac ctg ttt gaa cac atc aaa	576
Gly Ile Ser Ala Ser Ile Val Gly Ile Asp Leu Phe Glu His Ile Lys	
180 185 190	
aat atg atg tac acc acc att ccc gcg tgg ctc atc agt gcg gca ctg	624
Asn Met Met Tyr Thr Thr Ile Pro Ala Trp Leu Ile Ser Ala Ala Leu	
195 200 205	
atg ctg tgg ctt ctt ccc agc gtc gcc gcg cag gat ttg aac agc gtc	672
Met Leu Trp Leu Leu Pro Ser Val Ala Ala Gln Asp Leu Asn Ser Val	
210 215 220	
gaa tcc ttc cgc agc cag ctt gaa gcc acg gga ttg gtg cac tgc tat	720
Glu Ser Phe Arg Ser Gln Leu Glu Ala Thr Gly Leu Val His Cys Tyr	
225 230 235 240	
tcg ctg att ccg ttt gcg ctg ttg gtc gtt ttg gca ttg atg cgc gtc	768
Ser Leu Ile Pro Phe Ala Leu Leu Val Val Leu Ala Leu Met Arg Val	
245 250 255	
aat gcc gtg gtc gct atg ctc ttt acc gtc att gcc gcc gtt gcc gta	816
Asn Ala Val Val Ala Met Leu Phe Thr Val Ile Ala Ala Val Ala Val	
260 265 270	
acg tat ctg cac agc acg ccc gat ctg cgt cag ctc ggc gcg tgg ttt	864

Thr Tyr Leu His Ser Thr Pro Asp Leu Arg Gln Leu Gly Ala Trp Phe	
275 280 285	
tac ggc ggc tac aaa ctc gaa ggc gaa gcg ttt aaa gac att gcc aaa	912
Tyr Gly Gly Tyr Lys Leu Glu Gly Glu Ala Phe Lys Asp Ile Ala Lys	
290 295 300	
ctc att tct cgc ggc ggt ttg gaa agt atg ttt ttc acg cag acc atc	960
Leu Ile Ser Arg Gly Gly Leu Glu Ser Met Phe Phe Thr Gln Thr Ile	
305 310 315 320	
gtg att ctt ggg atg agc ctt ggc ggg ctg ctg ttt gca ctg ggc gcg	1008
Val Ile Leu Gly Met Ser Leu Gly Gly Leu Leu Phe Ala Leu Gly Ala	
325 330 335	
att cct tcc ctg ctg gat gcc gtc cgc agc ttt ttg acg aat gcc ggg	1056
Ile Pro Ser Leu Leu Asp Ala Val Arg Ser Phe Leu Thr Asn Ala Gly	
340 345 350	
cgt gcc aca ttc agc gtt gcc atg act tcg gtc ggg gtt aat ttc ctg	1104
Arg Ala Thr Phe Ser Val Ala Met Thr Ser Val Gly Val Asn Phe Leu	
355 360 365	
atc ggc gag caa tat ttg agt att ttg ttg tcg ggt gaa acg ttc aaa	1152
Ile Gly Glu Gln Tyr Leu Ser Ile Leu Leu Ser Gly Glu Thr Phe Lys	
370 375 380	
cct gtt tac gat aag ctc ggt ctg cat tcg cgc aat ctg tcg cgg acg	1200
Pro Val Tyr Asp Lys Leu Gly Leu His Ser Arg Asn Leu Ser Arg Thr	
385 390 395 400	
ctg gaa gat gcg ggg acg gtc atc aac ccg ctc gta ccg tgg agc gta	1248
Leu Glu Asp Ala Gly Thr Val Ile Asn Pro Leu Val Pro Trp Ser Val	
405 410 415	
tgc ggc gtg ttc atc agc cac gcg ctg ggc gtg ccg gtt tgg gaa tat	1296
Cys Gly Val Phe Ile Ser His Ala Leu Gly Val Pro Val Trp Glu Tyr	
420 425 430	
ctg ccg tat gcc ttt ttc tgc tat ttg agt ttg gct ttg acc ctg tta	1344
Leu Pro Tyr Ala Phe Phe Cys Tyr Leu Ser Leu Ala Leu Thr Leu Leu	
435 440 445	
ttc ggt tgg acg ggg ctg act ttg agc aaa aaa	1377
Phe Gly Trp Thr Gly Leu Thr Leu Ser Lys Lys	
450 455	

<210> 62

<211> 459

<212> PRT

<213> Neisseria meningitidis

<400> 62

Met Phe Ala Phe Lys Ser Leu Leu Asp Met Pro Arg Gly Glu Ala Leu
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Ala Val Val Val Ala Leu Ile Ala Ala Met Gly Tyr Thr Ile Ile Ser
 20 25 30

Leu Glu Trp Leu Pro His Met Ser Ile Ile Ala Ala Ile Val Val Leu
 35 40 45

Ile Leu Tyr Gly Leu Ala Arg Gly Leu Lys Tyr Asn Asp Met Gln Gln
 50 55 60

Gly Met Ile Gly Ala Leu Asn Gln Gly Met Gly Ala Ile Tyr Leu Phe
 65 70 75 80

Phe Phe Ile Gly Leu Met Val Ser Ala Leu Met Met Ser Gly Ala Ile
 85 90 95

Pro Thr Leu Met Tyr Tyr Gly Phe Gly Leu Ile Ser Pro Thr Tyr Phe
 100 105 110

Tyr Phe Ser Ala Phe Ala Leu Cys Ser Val Ile Gly Val Ser Ile Gly
 115 120 125

Ser Ser Leu Thr Thr Cys Ala Thr Val Gly Val Ala Phe Met Gly Met
 130 135 140

Ala Ala Ala Phe Gln Ala Asp Met Ala Met Thr Ala Gly Ala Ile Val
 145 150 155 160

Ser Gly Ala Phe Phe Gly Asp Lys Met Ser Pro Leu Ser Asp Thr Thr
 165 170 175

Gly Ile Ser Ala Ser Ile Val Gly Ile Asp Leu Phe Glu His Ile Lys
 180 185 190

Asn Met Met Tyr Thr Thr Ile Pro Ala Trp Leu Ile Ser Ala Ala Leu
 195 200 205

Met Leu Trp Leu Leu Pro Ser Val Ala Ala Gln Asp Leu Asn Ser Val
 210 215 220

Glu Ser Phe Arg Ser Gln Leu Glu Ala Thr Gly Leu Val His Cys Tyr
225 230 235 240

Ser Leu Ile Pro Phe Ala Leu Leu Val Val Leu Ala Leu Met Arg Val
245 250 255

Asn Ala Val Val Ala Met Leu Phe Thr Val Ile Ala Ala Val Ala Val
260 265 270

Thr Tyr Leu His Ser Thr Pro Asp Leu Arg Gln Leu Gly Ala Trp Phe
275 280 285

Tyr Gly Gly Tyr Lys Leu Glu Gly Glu Ala Phe Lys Asp Ile Ala Lys
290 295 300

Leu Ile Ser Arg Gly Gly Leu Glu Ser Met Phe Phe Thr Gln Thr Ile
305 310 315 320

Val Ile Leu Gly Met Ser Leu Gly Gly Leu Leu Phe Ala Leu Gly Ala
325 330 335

Ile Pro Ser Leu Leu Asp Ala Val Arg Ser Phe Leu Thr Asn Ala Gly
340 345 350

Arg Ala Thr Phe Ser Val Ala Met Thr Ser Val Gly Val Asn Phe Leu
355 360 365

Ile Gly Glu Gln Tyr Leu Ser Ile Leu Leu Ser Gly Glu Thr Phe Lys
370 375 380

Pro Val Tyr Asp Lys Leu Gly Leu His Ser Arg Asn Leu Ser Arg Thr
385 390 395 400

Leu Glu Asp Ala Gly Thr Val Ile Asn Pro Leu Val Pro Trp Ser Val
405 410 415

Cys Gly Val Phe Ile Ser His Ala Leu Gly Val Pro Val Trp Glu Tyr
420 425 430

Leu Pro Tyr Ala Phe Phe Cys Tyr Leu Ser Leu Ala Leu Thr Leu Leu
435 440 445

Phe Gly Trp Thr Gly Leu Thr Leu Ser Lys Lys
450 455

<211> 1098

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(1098)

<400> 63

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atg gca ggc aac act ttc gga caa ctc ttc acc gtt acc acc ttc ggc 48
Met Ala Gly Asn Thr Phe Gly Gln Leu Phe Thr Val Thr Thr Phe Gly
      1              5              10              15

gaa agc cac ggc gcg ggt ttg ggc tgt atc atc gac ggc tgc ccg ccc 96
Glu Ser His Gly Ala Gly Leu Gly Cys Ile Ile Asp Gly Cys Pro Pro
              20              25              30

ggc ttg gaa tta agc gaa gcg gat atc caa ttt gac ctc gac cga cgc 144
Gly Leu Glu Leu Ser Glu Ala Asp Ile Gln Phe Asp Leu Asp Arg Arg
      35              40              45

aaa ccc ggc acc agc cgc cac gtt acc caa cgc cgc gaa gcc gac caa 192
Lys Pro Gly Thr Ser Arg His Val Thr Gln Arg Arg Glu Ala Asp Gln
      50              55              60

gtc gaa atc ctc tcc ggc gta ttc gaa ggc aaa acc acc ggc acg ccc 240
Val Glu Ile Leu Ser Gly Val Phe Glu Gly Lys Thr Thr Gly Thr Pro
      65              70              75              80

atc gcc ctc tta atc cgc aat acc gac cag cgc agc aaa gac tac ggc 288
Ile Ala Leu Leu Ile Arg Asn Thr Asp Gln Arg Ser Lys Asp Tyr Gly
              85              90              95

aac atc gcc acc agc ttc cgc ccc ggc cac gcc gac tat acc tat tgg 336
Asn Ile Ala Thr Ser Phe Arg Pro Gly His Ala Asp Tyr Thr Tyr Trp
              100              105              110

cac aaa tac ggc acg cgc gac tac cgg ggc ggc ggc agg agt tcc gcc 384
His Lys Tyr Gly Thr Arg Asp Tyr Arg Gly Gly Gly Arg Ser Ser Ala
              115              120              125

cgc gaa acc gcc gcc cgc gtt gcc gcc gga gcc gtt gcc aaa aaa tgg 432
Arg Glu Thr Ala Ala Arg Val Ala Ala Gly Ala Val Ala Lys Lys Trp
              130              135              140

ttg aaa gaa aaa ttc ggc acg gaa atc acc gcc tac gtt acc caa gtc 480
Ile Lys Glu Lys Phe Gly Thr Glu Ile Thr Ala Tyr Val Thr Gln Val
      145              150              155              160

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ggc gaa aaa gaa atc cgg ttt gaa ggc tgc gaa cac att tcc caa aat 528
Gly Glu Lys Glu Ile Arg Phe Glu Gly Cys Glu His Ile Ser Gln Asn
      165              170              175

cct ttt ttt gcc gcc aac cat agc caa att gcc gag ctg gaa aac tat 576
Pro Phe Phe Ala Ala Asn His Ser Gln Ile Ala Glu Leu Glu Asn Tyr
      180              185              190

atg gac agc gtg cgc aaa tcc ttg gat tcc gtc ggc gcg aag ctg cat 624
Met Asp Ser Val Arg Lys Ser Leu Asp Ser Val Gly Ala Lys Leu His
      195              200              205

att gaa gcc gcc aac gtc ccc gtc ggc ttg ggc gaa ccc gta ttc gac 672
Ile Glu Ala Ala Asn Val Pro Val Gly Leu Gly Glu Pro Val Phe Asp
      210              215              220

cgc ctc gat gcc gaa atc gcc tac gcc atg atg ggc atc aac gcc gtc 720
Arg Leu Asp Ala Glu Ile Ala Tyr Ala Met Met Gly Ile Asn Ala Val
      225              230              235              240

aaa ggc gtg gaa atc ggc gca ggt ttt gac agc gta acg caa cgc ggc 768
Lys Gly Val Glu Ile Gly Ala Gly Phe Asp Ser Val Thr Gln Arg Gly
      245              250              255

agc gaa cac ggc gac gaa ctc acc ccg caa ggc ttc ctg tcc aac cac 816
Ser Glu His Gly Asp Glu Leu Thr Pro Gln Gly Phe Leu Ser Asn His
      260              265              270

tca ggc ggc atc ctc ggc ggc atc agc acc gga caa gac atc cac gtc 864
Ser Gly Gly Ile Leu Gly Gly Ile Ser Thr Gly Gln Asp Ile His Val
      275              280              285

aat atc gcc atc aaa ccc acc agc tcc atc gcc acc ccg cgc cgc agt 912
Asn Ile Ala Ile Lys Pro Thr Ser Ser Ile Ala Thr Pro Arg Arg Ser
      290              295              300

atc gac atc aac ggc aac ccc atc gaa ctc gcc acg cac ggc agg cac 960
Ile Asp Ile Asn Gly Asn Pro Ile Glu Leu Ala Thr His Gly Arg His
      305              310              315              320

gac ccc tgc gtc ggc ttg cgc gcc gcg ccg att gca gaa gct atg ctg 1008
Asp Pro Cys Val Gly Leu Arg Ala Ala Pro Ile Ala Glu Ala Met Leu
      325              330              335

gca ttg gtt tta atc gac cac gcc ctg cgc cat cgc gcg caa aat gcg 1056
Ala Leu Val Leu Ile Asp His Ala Leu Arg His Arg Ala Gln Asn Ala
      340              345              350

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gat gtt cag gtt aat acg ccc gac att acc ctt tca aac aaa 1098
 Asp Val Gln Val Asn Thr Pro Asp Ile Thr Leu Ser Asn Lys
 355 360 365

<210> 64

<211> 366

<212> PRT

<213> Neisseria meningitidis

<400> 64

Met Ala Gly Asn Thr Phe Gly Gln Leu Phe Thr Val Thr Thr Phe Gly
 1 5 10 15

Glu Ser His Gly Ala Gly Leu Gly Cys Ile Ile Asp Gly Cys Pro Pro
 20 25 30

Gly Leu Glu Leu Ser Glu Ala Asp Ile Gln Phe Asp Leu Asp Arg Arg
 35 40 45

Lys Pro Gly Thr Ser Arg His Val Thr Gln Arg Arg Glu Ala Asp Gln
 50 55 60

Val Glu Ile Leu Ser Gly Val Phe Glu Gly Lys Thr Thr Gly Thr Pro
 65 70 75 80

Ile Ala Leu Leu Ile Arg Asn Thr Asp Gln Arg Ser Lys Asp Tyr Gly
 85 90 95

Asn Ile Ala Thr Ser Phe Arg Pro Gly His Ala Asp Tyr Thr Tyr Trp
 100 105 110

His Lys Tyr Gly Thr Arg Asp Tyr Arg Gly Gly Gly Arg Ser Ser Ala
 115 120 125

Arg Glu Thr Ala Ala Arg Val Ala Ala Gly Ala Val Ala Lys Lys Trp
 130 135 140

Leu Lys Glu Lys Phe Gly Thr Glu Ile Thr Ala Tyr Val Thr Gln Val
 145 150 155 160

Gly Glu Lys Glu Ile Arg Phe Glu Gly Cys Glu His Ile Ser Gln Asn
 165 170 175

Pro Phe Phe Ala Ala Asn His Ser Gln Ile Ala Glu Leu Glu Asn Tyr
 180 185 190

Met Asp Ser Val Arg Lys Ser Leu Asp Ser Val Gly Ala Lys Leu His
 195 200 205

Ile Glu Ala Ala Asn Val Pro Val Gly Leu Gly Glu Pro Val Phe Asp
 210 215 220

Arg Leu Asp Ala Glu Ile Ala Tyr Ala Met Met Gly Ile Asn Ala Val
 225 230 235 240

Lys Gly Val Glu Ile Gly Ala Gly Phe Asp Ser Val Thr Gln Arg Gly
 245 250 255

Ser Glu His Gly Asp Glu Leu Thr Pro Gln Gly Phe Leu Ser Asn His
 260 265 270

Ser Gly Gly Ile Leu Gly Gly Ile Ser Thr Gly Gln Asp Ile His Val
 275 280 285

Asn Ile Ala Ile Lys Pro Thr Ser Ser Ile Ala Thr Pro Arg Arg Ser
 290 295 300

Ile Asp Ile Asn Gly Asn Pro Ile Glu Leu Ala Thr His Gly Arg His
 305 310 315 320

Asp Pro Cys Val Gly Leu Arg Ala Ala Pro Ile Ala Glu Ala Met Leu
 325 330 335

Ala Leu Val Leu Ile Asp His Ala Leu Arg His Arg Ala Gln Asn Ala
 340 345 350

Asp Val Gln Val Asn Thr Pro Asp Ile Thr Leu Ser Asn Lys
 355 360 365

<210> 65

<211> 461

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(459)

<400> 65

atg caa agc ggt ttt aac gct att ttc agg aat gac acc atg cag gtt 48
 Met Gln Ser Gly Phe Asn Ala Ile Phe Arg Asn Asp Thr Met Gln Val
 1 5 10 15

aca tca aaa tgg ata gac gga atg tgc ttt gtc gga aca acg gaa ggc 96
 Thr Ser Lys Trp Ile Asp Gly Met Cys Phe Val Gly Thr Thr Glu Gly
 20 25 30

ggg cac agc gtc gtt atg gag ggg tcg gcg gca gaa ggt gcg gcc aag 144
 Gly His Ser Val Val Met Glu Gly Ser Ala Ala Glu Gly Ala Ala Lys
 35 40 45

cgc ggg ccc agc cct ttg gaa atg ctg ctg ttg ggc gtg gcg ggc tgt 192
 Arg Gly Pro Ser Pro Leu Glu Met Leu Leu Leu Gly Val Ala Gly Cys
 50 55 60

tcg agc atc gat gtg gtg atg att gcc gaa aaa cag cgt cag aaa gtg 240
 Ser Ser Ile Asp Val Val Met Ile Ala Glu Lys Gln Arg Gln Lys Val
 65 70 75 80

act gac tgc cgt gcg acg gtt acg gcg aaa cgg gcg gac gat gcg ccg 288
 Thr Asp Cys Arg Ala Thr Val Thr Ala Lys Arg Ala Asp Asp Ala Pro
 85 90 95

cgc gtg ttt acc gaa atc cac atc cat ttc aaa gta ttc ggg cat gat 336
 Arg Val Phe Thr Glu Ile His Ile His Phe Lys Val Phe Gly His Asp
 100 105 110

ttg aaa gaa tcg gcc att gag cgc gcc gtt cag atg tct gcc gaa aaa 384
 Leu Lys Glu Ser Ala Ile Glu Arg Ala Val Gln Met Ser Ala Glu Lys
 115 120 125

tac tgt tcg gct tcg att atg ttg ggc aaa gcg gca aag att acc cac 432
 Tyr Cys Ser Ala Ser Ile Met Leu Gly Lys Ala Ala Lys Ile Thr His
 130 135 140

agt ttt gaa att gcc ggg gca gat aaa ta 461
 Ser Phe Glu Ile Ala Gly Ala Asp Lys
 145 150

<210> 66

<211> 153

<212> PRT

<213> Neisseria meningitidis

<400> 66

Met Gln Ser Gly Phe Asn Ala Ile Phe Arg Asn Asp Thr Met Gln Val
 1 5 10 15

Thr Ser Lys Trp Ile Asp Gly Met Cys Phe Val Gly Thr Thr Glu Gly

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<210> 67
<211> 1659
<212> DNA
<213> Neisseria meningitidis
<220>
<221> CDS
<222> {1}..{1659}
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<400> 67																	
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Met	Thr	Asp	Asn	Ser	Pro	Pro	Pro.	Asn	Gly	His	Ala	Gln	Ala	Arg	Val		
1				5					10					15			
cgc	aaa	aac	aac	acc	ttc	ctc	tct	gcc	gtc	tgg	ctg	gtt	ccg	ctg	atc	96	
Arg	Lys	Asn	Asn	Thr	Phe	Leu	Ser	Ala	Val	Trp	Leu	Val	Pro	Leu	Ile		
				20					25					30			
gcg	ctg	att	gcc	ggc	ggc	tgg	ctt	tgg	gtt	aag	gaa	atc	cgc	aac	agg	144	
Ala	Leu	Ile	Ala	Gly	Gly	Trp	Leu	Trp	Val	Lys	Glu	Ile	Arg	Asn	Arg		

35	40	45	
ggg cct gtg gtt acg ctc ttg atg gac agc gcg gaa ggc att gag gtc			192
Gly Pro Val Val Thr Leu Leu Met Asp Ser Ala Glu Gly Ile Glu Val			
50	55	60	
aac aat acg gtc atc aaa gta ttg agc atc gat gtc gga cgc gtt acc			240
Asn Asn Thr Val Ile Lys Val Leu Ser Ile Asp Val Gly Arg Val Thr			
65	70	75	80
cga atc aaa ctg cgc gac gac caa aaa ggc gtg gaa gta acc gcc caa			288
Arg Ile Lys Leu Arg Asp Asp Gln Lys Gly Val Glu Val Thr Ala Gln			
85	90	95	
ctc aat gcg gac gta tcc ggc ctc atc cgc agc gat acc cag ttt tgg			336
Leu Asn Ala Asp Val Ser Gly Leu Ile Arg Ser Asp Thr Gln Phe Trp			
100	105	110	
gtg gtc aag ccg cgt atc gac caa agc ggc gta acc ggt ttg ggt acg			384
Val Val Lys Pro Arg Ile Asp Gln Ser Gly Val Thr Gly Leu Gly Thr			
115	120	125	
ctg ctt tcg ggt tcg tac atc gcc ttt aca ccc ggc aaa agc gac gag			432
Leu Leu Ser Gly Ser Tyr Ile Ala Phe Thr Pro Gly Lys Ser Asp Glu			
130	135	140	
gca aaa gac gtg ttc caa gtg cag gac att ccg ccc gtt acc gcc atc			480
Ala Lys Asp Val Phe Gln Val Gln Asp Ile Pro Pro Val Thr Ala Ile			
145	150	155	160
ggg caa agc ggg ctg cgc ttg aat ttg att ggt aaa aac gac cgc atc			528
Gly Gln Ser Gly Leu Arg Leu Asn Leu Ile Gly Lys Asn Asp Arg Ile			
165	170	175	
ctc aac gtc aac agc cct gtt ttg tat gaa aac ttt atg gtc ggg caa			576
Leu Asn Val Asn Ser Pro Val Leu Tyr Glu Asn Phe Met Val Gly Gln			
180	185	190	
gtc gaa agc gcg cat ttc gac ccg tcc gac caa agc gtg cat tac acc			624
Val Glu Ser Ala His Phe Asp Pro Ser Asp Gln Ser Val His Tyr Thr			
195	200	205	
atc ttc atc caa agc ccc aac gac aaa ctg att cat tcc gcc agc cgt			672
Ile Phe Ile Gln Ser Pro Asn Asp Lys Leu Ile His Ser Ala Ser Arg			
210	215	220	
ttc tgg ctg gaa agc ggc atc aat atc gaa acc aca ggc agc ggc atc			720
Phe Trp Leu Glu Ser Gly Ile Asn Ile Glu Thr Thr Gly Ser Gly Ile			

225	230	235	240	
aaa ctc aat tcc gcc cct ctg cct gcc ctg ctg tcg ggc gcg att tca				768
Lys Leu Asn Ser Ala Pro Leu Pro Ala Leu Leu Ser Gly Ala Ile Ser				
	245	250	255	
ttt gat tcg ccg aaa acc aaa aac agt aaa aac gtc aaa agc gaa gac				816
Phe Asp Ser Pro Lys Thr Lys Asn Ser Lys Asn Val Lys Ser Glu Asp				
	260	265	270	
agc ttc acg ctt tac gac agc cgc agc gaa gtc gcc aac ctg cct gat				864
Ser Phe Thr Leu Tyr Asp Ser Arg Ser Glu Val Ala Asn Leu Pro Asp				
	275	280	285	
gac cgt tcg ctg tac tac acc gcg ttt ttc aaa caa tcc gtg cgc gga				912
Asp Arg Ser Leu Tyr Tyr Thr Ala Phe Phe Lys Gln Ser Val Arg Gly				
	290	295	300	
ctg acc gtc ggt tcg cct gtc gag tac aaa ggg ctg aat gtc ggc gtg				960
Leu Thr Val Gly Ser Pro Val Glu Tyr Lys Gly Leu Asn Val Gly Val				
305	310	315	320	
gtt tcc gat gtt cct tat ttc gac cgc aac gac agc ctg cac ctg ttt				1008
Val Ser Asp Val Pro Tyr Phe Asp Arg Asn Asp Ser Leu His Leu Phe				
	325	330	335	
gaa aac ggc tgg att ccc gta cgc atc cgt att gag cct tcc cgt ttg				1056
Glu Asn Gly Trp Ile Pro Val Arg Ile Arg Ile Glu Pro Ser Arg Leu				
	340	345	350	
gaa atc aat gcc gac gaa caa agc aaa gaa cat tgg aaa caa caa ttt				1104
Glu Ile Asn Ala Asp Glu Gln Ser Lys Glu His Trp Lys Gln Gln Phe				
	355	360	365	
cag acg gcc tta aac aaa ggc ctg acc gcc acc atc tcc agc aac aac				1152
Gln Thr Ala Leu Asn Lys Gly Leu Thr Ala Thr Ile Ser Ser Asn Asn				
	370	375	380	
ctg ctg acc ggc agc aaa atg att gag ttg aac gat cag cct tcc gcc				1200
Leu Leu Thr Gly Ser Lys Met Ile Glu Leu Asn Asp Gln Pro Ser Ala				
385	390	395	400	
tcg ccc aag ctg cga ccg cat acc gtt tat gca ggc gat acc gtt atc				1248
Ser Pro Lys Leu Arg Pro His Thr Val Tyr Ala Gly Asp Thr Val Ile				
	405	410	415	
gcg acc cag ggc ggc ggt ttg gac gat ttg cag gtc aaa ttg gcg gat				1296
Ala Thr Gln Gly Gly Gly Leu Asp Asp Leu Gln Val Lys Leu Ala Asp				

420	425	430	
ttg ctg gac aag ttc gac aaa ctg cct tta gat aag acg gtt gcc gaa			1344
Leu Leu Asp Lys Phe Asp Lys Leu Pro Leu Asp Lys Thr Val Ala Glu			
435	440	445	
ttg aac ggt tcg ctt gcc gag ctc aaa tcc aca ctc aaa tct gcc aat			1392
Leu Asn Gly Ser Leu Ala Glu Leu Lys Ser Thr Leu Lys Ser Ala Asn			
450	455	460	
gcc gcc cta agc tcc atc gac aaa ctg gtc ggc aaa ccg cag aca caa			1440
Ala Ala Leu Ser Ser Ile Asp Lys Leu Val Gly Lys Pro Gln Thr Gln			
465	470	475	480
aac att ccg aac gaa ctg aac caa acc ctg aaa gag ttg cgc aca acc			1488
Asn Ile Pro Asn Glu Leu Asn Gln Thr Leu Lys Glu Leu Arg Thr Thr			
485	490	495	
ctg caa ggc gta tcg cct caa tcg cct atc tac ggc gac gta caa aat			1536
Leu Gln Gly Val Ser Pro Gln Ser Pro Ile Tyr Gly Asp Val Gln Asn			
500	505	510	
acg ctg caa agt ttg gac aaa acc tta aaa gac gtt caa ccc gtc att			1584
Thr Leu Gln Ser Leu Asp Lys Thr Leu Lys Asp Val Gln Pro Val Ile			
515	520	525	
aac act ttg aaa gaa aaa ccc aac gcg ctg att ttc aac agc agc agc			1632
Asn Thr Leu Lys Glu Lys Pro Asn Ala Leu Ile Phe Asn Ser Ser Ser			
530	535	540	
aaa gac cct atc ccg aaa gga agc cga			1659
Lys Asp Pro Ile Pro Lys Gly Ser Arg			
545	550		

<210> 68

<211> 553

<212> PRT

<213> Neisseria meningitidis

<400> 68

Met	Thr	Asp	Asn	Ser	Pro	Pro	Pro	Asn	Gly	His	Ala	Gln	Ala	Arg	Val
1				5					10					15	

Arg	Lys	Asn	Asn	Thr	Phe	Leu	Ser	Ala	Val	Trp	Leu	Val	Pro	Leu	Ile
				20					25					30	

Ala	Leu	Ile	Ala	Gly	Gly	Trp	Leu	Trp	Val	Lys	Glu	Ile	Arg	Asn	Arg
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

35	40	45
Gly Pro Val Val Thr Leu Leu Met Asp Ser Ala Glu Gly Ile Glu Val		
50	55	60
Asn Asn Thr Val Ile Lys Val Leu Ser Ile Asp Val Gly Arg Val Thr		
65	70	75
Arg Ile Lys Leu Arg Asp Asp Gln Lys Gly Val Glu Val Thr Ala Gln		
85	90	95
Leu Asn Ala Asp Val Ser Gly Leu Ile Arg Ser Asp Thr Gln Phe Trp		
100	105	110
Val Val Lys Pro Arg Ile Asp Gln Ser Gly Val Thr Gly Leu Gly Thr		
115	120	125
Leu Leu Ser Gly Ser Tyr Ile Ala Phe Thr Pro Gly Lys Ser Asp Glu		
130	135	140
Ala Lys Asp Val Phe Gln Val Gln Asp Ile Pro Pro Val Thr Ala Ile		
145	150	155
Gly Gln Ser Gly Leu Arg Leu Asn Leu Ile Gly Lys Asn Asp Arg Ile		
165	170	175
Leu Asn Val Asn Ser Pro Val Leu Tyr Glu Asn Phe Met Val Gly Gln		
180	185	190
Val Glu Ser Ala His Phe Asp Pro Ser Asp Gln Ser Val His Tyr Thr		
195	200	205
Ile Phe Ile Gln Ser Pro Asn Asp Lys Leu Ile His Ser Ala Ser Arg		
210	215	220
Phe Trp Leu Glu Ser Gly Ile Asn Ile Glu Thr Thr Gly Ser Gly Ile		
225	230	235
Lys Leu Asn Ser Ala Pro Leu Pro Ala Leu Leu Ser Gly Ala Ile Ser		
245	250	255
Phe Asp Ser Pro Lys Thr Lys Asn Ser Lys Asn Val Lys Ser Glu Asp		
260	265	270
Ser Phe Thr Leu Tyr Asp Ser Arg Ser Glu Val Ala Asn Leu Pro Asp		
275	280	285
Asp Arg Ser Leu Tyr Tyr Thr Ala Phe Phe Lys Gln Ser Val Arg Gly		

290	295	300	
Leu Thr Val Gly Ser Pro Val Glu Tyr Lys Gly Leu Asn Val Gly Val			
305	310	315	320
Val Ser Asp Val Pro Tyr Phe Asp Arg Asn Asp Ser Leu His Leu Phe			
	325	330	335
Glu Asn Gly Trp Ile Pro Val Arg Ile Arg Ile Glu Pro Ser Arg Leu			
	340	345	350
Glu Ile Asn Ala Asp Glu Gln Ser Lys Glu His Trp Lys Gln Gln Phe			
	355	360	365
Gln Thr Ala Leu Asn Lys Gly Leu Thr Ala Thr Ile Ser Ser Asn Asn			
	370	375	380
Leu Leu Thr Gly Ser Lys Met Ile Glu Leu Asn Asp Gln Pro Ser Ala			
385	390	395	400
Ser Pro Lys Leu Arg Pro His Thr Val Tyr Ala Gly Asp Thr Val Ile			
	405	410	415
Ala Thr Gln Gly Gly Gly Leu Asp Asp Leu Gln Val Lys Leu Ala Asp			
	420	425	430
Leu Leu Asp Lys Phe Asp Lys Leu Pro Leu Asp Lys Thr Val Ala Glu			
	435	440	445
Leu Asn Gly Ser Leu Ala Glu Leu Lys Ser Thr Leu Lys Ser Ala Asn			
	450	455	460
Ala Ala Leu Ser Ser Ile Asp Lys Leu Val Gly Lys Pro Gln Thr Gln			
465	470	475	480
Asn Ile Pro Asn Glu Leu Asn Gln Thr Leu Lys Glu Leu Arg Thr Thr			
	485	490	495
Leu Gln Gly Val Ser Pro Gln Ser Pro Ile Tyr Gly Asp Val Gln Asn			
	500	505	510
Thr Leu Gln Ser Leu Asp Lys Thr Leu Lys Asp Val Gln Pro Val Ile			
	515	520	525
Asn Thr Leu Lys Glu Lys Pro Asn Ala Leu Ile Phe Asn Ser Ser Ser			
	530	535	540
Lys Asp Pro Ile Pro Lys Gly Ser Arg			

545

550

<210> 69

<211> 330

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(330)

<400> 69

ttg	ccg	ctg	ctt	tgt	tgt	ctt	tgg	ttt	tgg	cag	cct	gcg	gcg	gtg	aaa	48
Leu	Pro	Leu	Leu	Cys	Cys	Leu	Trp	Phe	Trp	Gln	Pro	Ala	Ala	Val	Lys	
1				5					10					15		

aag	ccg	ctg	aag	ctc	ccg	ctg	ctg	aag	cac	ctg	ccg	ccg	aag	ctc	ccg	96
Lys	Pro	Leu	Lys	Leu	Pro	Leu	Leu	Lys	His	Leu	Pro	Pro	Lys	Leu	Pro	
			20					25					30			

cta	ctg	aag	cac	ctg	ccg	ccg	aag	ctc	ccg	ctg	ctg	aag	cac	ctg	ccg	144
Leu	Leu	Lys	His	Leu	Pro	Pro	Lys	Leu	Pro	Leu	Leu	Lys	His	Leu	Pro	
		35					40					45				

ccg	aag	ctc	ctg	cta	ctg	aag	cac	ctg	ccg	ccg	aag	ctc	ccg	ctg	ctg	192
Pro	Lys	Leu	Leu	Leu	Leu	Lys	His	Leu	Pro	Pro	Lys	Leu	Pro	Leu	Leu	
	50					55					60					

aag	ctg	ccg	cta	ccg	aag	cac	ctg	ccg	ctg	aag	ctg	ccg	cta	ccg	aag	240
Lys	Leu	Pro	Leu	Pro	Lys	His	Leu	Pro	Leu	Lys	Leu	Pro	Leu	Pro	Lys	
65					70				75					80		

cac	ctg	ccg	ctg	aag	ctc	ctg	ctg	ccg	aag	ctg	caa	aat	aag	cat	ttt	288
His	Leu	Pro	Leu	Lys	Leu	Leu	Leu	Pro	Lys	Leu	Gln	Asn	Lys	His	Phe	
			85					90					95			

ccg	ctt	gca	aaa	aag	cag	gat	acg	ttc	agt	atc	ctg	ctt	ttt			330
Pro	Leu	Ala	Lys	Lys	Gln	Asp	Thr	Phe	Ser	Ile	Leu	Leu	Phe			
		100					105					110				

<210> 70

<211> 110

<212> PRT

<213> Neisseria meningitidis

<400> 70

Leu Pro Leu Leu Cys Cys Leu Trp Phe Trp Gln Pro Ala Ala Val Lys
 1 5 10 15

Lys Pro Leu Lys Leu Pro Leu Leu Lys His Leu Pro Pro Lys Leu Pro
 20 25 30

Leu Leu Lys His Leu Pro Pro Lys Leu Pro Leu Leu Lys His Leu Pro
 35 40 45

Pro Lys Leu Leu Leu Leu Lys His Leu Pro Pro Lys Leu Pro Leu Leu
 50 55 60

Lys Leu Pro Leu Pro Lys His Leu Pro Leu Lys Leu Pro Leu Pro Lys
 65 70 75 80

His Leu Pro Leu Lys Leu Leu Leu Pro Lys Leu Gln Asn Lys His Phe
 85 90 95

Pro Leu Ala Lys Lys Gln Asp Thr Phe Ser Ile Leu Leu Phe
 100 105 110

<210> 71

<211> 2274

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(2274)

<400> 71

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 Met Thr Thr Leu His Phe Ser Gly Phe Pro Arg Val Gly Ala Phe Arg
 1 5 10 15

gaa ttg aaa ttc gca caa gaa aaa tac tgg cgc aaa gaa atc agc gag 96
 Glu Leu Lys Phe Ala Gln Glu Lys Tyr Trp Arg Lys Glu Ile Ser Glu
 20 25 30

caa gaa ttg ctg gct gtt gct aaa gac ttg cgc gag aaa aac tgg aaa 144
 Gln Glu Leu Leu Ala Val Ala Lys Asp Leu Arg Glu Lys Asn Trp Lys
 35 40 45

cac cag gcc gct gcc aac gcc gat tac gtt gcc gta ggc gat ttc act 192
 His Gln Ala Ala Ala Asn Ala Asp Tyr Val Ala Val Gly Asp Phe Thr

50	55	60	
ttc tac gac cac atc ctc gac ctg caa gtc gcc acc ggc gcg att ccc			240
Phe Tyr Asp His Ile Leu Asp Leu Gln Val Ala Thr Gly Ala Ile Pro			
65	70	75	80
gcc cgc ttc ggc ttc gac agc caa aac cta tct ttg gaa caa ttc ttc			288
Ala Arg Phe Gly Phe Asp Ser Gln Asn Leu Ser Leu Glu Gln Phe Phe			
	85	90	95
caa ctg gcg cgc ggt aac aaa gac caa ttc gct atc gaa atg acc aaa			336
Gln Leu Ala Arg Gly Asn Lys Asp Gln Phe Ala Ile Glu Met Thr Lys			
	100	105	110
tgg ttc gac acc aac tac cac tac ttg gtg cct gaa ttc cac gcc gat			384
Trp Phe Asp Thr Asn Tyr His Tyr Leu Val Pro Glu Phe His Ala Asp			
	115	120	125
acc gaa ttt aaa gcc aat gcc aaa cac tac gtt caa cag ctg caa gaa			432
Thr Glu Phe Lys Ala Asn Ala Lys His Tyr Val Gln Gln Leu Gln Glu			
	130	135	140
gcc caa gct ttg ggc ctg aaa gcc aaa ccg acc gtt gtg ggt ccg ttg			480
Ala Gln Ala Leu Gly Leu Lys Ala Lys Pro Thr Val Val Gly Pro Leu			
	145	150	155
act ttc ctg tgg gtc ggt aaa gaa aaa ggc gcc gtt gaa ttc gac cgt			528
Thr Phe Leu Trp Val Gly Lys Glu Lys Gly Ala Val Glu Phe Asp Arg			
	165	170	175
ctg agc ctg ttg cct aaa ctg ttg cct gtt tac gtt gaa atc ctg act			576
Leu Ser Leu Leu Pro Lys Leu Leu Pro Val Tyr Val Glu Ile Leu Thr			
	180	185	190
gct ttg gtt gaa gcc ggt gcc gag tgg att caa atc gac gag cct gct			624
Ala Leu Val Glu Ala Gly Ala Glu Trp Ile Gln Ile Asp Glu Pro Ala			
	195	200	205
ttg act gtt gac ctg cct aaa gaa tgg gta gaa gca tac aaa gac gtt			672
Leu Thr Val Asp Leu Pro Lys Glu Trp Val Glu Ala Tyr Lys Asp Val			
	210	215	220
tac gcc act ttg agc aaa gta agt gcc aaa atc ctg ctg agc act tac			720
Tyr Ala Thr Leu Ser Lys Val Ser Ala Lys Ile Leu Leu Ser Thr Tyr			
	225	230	235
ttc ggt tct gtt gcc gaa cac gcc gca ttg ttg aaa tcc ctg cct gtt			768
Phe Gly Ser Val Ala Glu His Ala Ala Leu Leu Lys Ser Leu Pro Val			

245	250	255	
gac ggc ctg cac atc gac ttg gta cgc gct cct gaa caa ctg gat gcg			816
Asp Gly Leu His Ile Asp Leu Val Arg Ala Pro Glu Gln Leu Asp Ala			
260	265	270	
ttc gcc gac tac gac aaa gtt ctg tct gcc ggc gtc att gac ggc cgc			864
Phe Ala Asp Tyr Asp Lys Val Leu Ser Ala Gly Val Ile Asp Gly Arg			
275	280	285	
aat att tgg cgc gcc aac ctg aac aaa gtt ttg gaa act gtt gag ctt			912
Asn Ile Trp Arg Ala Asn Leu Asn Lys Val Leu Glu Thr Val Glu Leu			
290	295	300	
ctg caa gcc aaa ctg ggc gac cgt ttg tgg att tcc agc tct tgc tcg			960
Leu Gln Ala Lys Leu Gly Asp Arg Leu Trp Ile Ser Ser Ser Cys Ser			
305	310	315	320
ctg ctg cac act ccg ttt gac ttg tca gtt gaa gaa aaa ctg aaa gcc			1008
Leu Leu His Thr Pro Phe Asp Leu Ser Val Glu Glu Lys Leu Lys Ala			
325	330	335	
aac aaa ccc gac ctg tac tct tgg ttg gca ttc acc ctg caa aaa acc			1056
Asn Lys Pro Asp Leu Tyr Ser Trp Leu Ala Phe Thr Leu Gln Lys Thr			
340	345	350	
caa gaa ttg cgc gtc ctg aaa gcc gca ttg aac gaa ggc cgt gat tct			1104
Gln Glu Leu Arg Val Leu Lys Ala Ala Leu Asn Glu Gly Arg Asp Ser			
355	360	365	
gtt gcc gaa gaa ctg gct gcc agc caa gct gct gcc gac tcc cgt gcc			1152
Val Ala Glu Glu Leu Ala Ala Ser Gln Ala Ala Ala Asp Ser Arg Ala			
370	375	380	
aac agc agc gaa atc cac cgt gca gac gtt gcc aaa cgc ctg gcc gat			1200
Asn Ser Ser Glu Ile His Arg Ala Asp Val Ala Lys Arg Leu Ala Asp			
385	390	395	400
ttg cct gcc aac gca gac caa cgc aaa tcc cca ttt gcc gac cgt atc			1248
Leu Pro Ala Asn Ala Asp Gln Arg Lys Ser Pro Phe Ala Asp Arg Ile			
405	410	415	
aaa gcg caa caa gca tgg ttg aac ctg cct ctg ctg ccg acg acc aac			1296
Lys Ala Gln Gln Ala Trp Leu Asn Leu Pro Leu Leu Pro Thr Thr Asn			
420	425	430	
atc ggt tct ttc ccg caa act acc gaa atc cgc caa gca cgc gca gcc			1344
Ile Gly Ser Phe Pro Gln Thr Thr Glu Ile Arg Gln Ala Arg Ala Ala			

435	440	445	
ttc aaa aaa ggc gaa ctg tct gcc gcc gat tac gaa gcc gcg atg aaa			1392
Phe Lys Lys Gly Glu Leu Ser Ala Ala Asp Tyr Glu Ala Ala Met Lys			
450	455	460	
aaa gaa atc gcc ttg gtg gtt gaa gag caa gaa aaa ctg gac ttg gac			1440
Lys Glu Ile Ala Leu Val Val Glu Glu Gln Glu Lys Leu Asp Leu Asp			
465	470	475	480
gta ctg gta cac ggc gaa gcc gag cgt aac gac atg gtt gaa tac ttc			1488
Val Leu Val Val His Gly Glu Ala Glu Arg Asn Asp Met Val Glu Tyr Phe			
485	490	495	
ggc gaa ttg ttg agc ggt ttt gca ttc acc caa tat ggc tgg gta caa			1536
Gly Glu Leu Leu Ser Gly Phe Ala Phe Thr Gln Tyr Gly Trp Val Gln			
500	505	510	
agc tac ggt tct cgc tgc gtt aaa cca ccg att atc ttc ggt gac gta			1584
Ser Tyr Gly Ser Arg Cys Val Lys Pro Pro Ile Ile Phe Gly Asp Val			
515	520	525	
agc cgt cct gaa gcc atg acc gta gct tgg tct act tac gca caa agc			1632
Ser Arg Pro Glu Ala Met Thr Val Ala Trp Ser Thr Tyr Ala Gln Ser			
530	535	540	
ctg acc aaa cgc ccg atg aaa ggt atg ttg acc ggc cct gta acc att			1680
Leu Thr Lys Arg Pro Met Lys Gly Met Leu Thr Gly Pro Val Thr Ile			
545	550	555	560
ctg caa tgg tct ttc gtc cgc aac gat att cct cgc tct acc gtg tgt			1728
Leu Gln Trp Ser Phe Val Arg Asn Asp Ile Pro Arg Ser Thr Val Cys			
565	570	575	
aaa caa atc gca ctg gct ctg aac gac gaa gta ttg gat ctg gaa aaa			1776
Lys Gln Ile Ala Leu Ala Leu Asn Asp Glu Val Leu Asp Leu Glu Lys			
580	585	590	
gca ggc atc aaa gtc atc caa atc gac gaa cct gcc atc cgc gaa ggc			1824
Ala Gly Ile Lys Val Ile Gln Ile Asp Glu Pro Ala Ile Arg Glu Gly			
595	600	605	
ctg cct ttg aaa cgt gcc gat tgg gat gcc tac ctg aac tgg gcc ggc			1872
Leu Pro Leu Lys Arg Ala Asp Trp Asp Ala Tyr Leu Asn Trp Ala Gly			
610	615	620	
gaa tct ttc cgc ctg tcc tct acc ggt tgc gaa gac agc act caa atc			1920
Glu Ser Phe Arg Leu Ser Ser Thr Gly Cys Glu Asp Ser Thr Gln Ile			

625	630	635	640	
cac acc cat atg tgc tac tct gag ttc aac gac atc ctg cct gcg att				1968
His Thr His Met Cys Tyr Ser Glu Phe Asn Asp Ile Leu Pro Ala Ile				
	645	650	655	
gct gca atg gat gcg gac gtg atc acc atc gag act tca cgt tcc gac				2016
Ala Ala Met Asp Ala Asp Val Ile Thr Ile Glu Thr Ser Arg Ser Asp				
	660	665	670	
atg gaa ctc ttg acc gcg ttc ggc gaa ttc aaa tac ccg aac gac atc				2064
Met Glu Leu Leu Thr Ala Phe Gly Glu Phe Lys Tyr Pro Asn Asp Ile				
	675	680	685	
ggc ccg ggg gtt tac gac atc cac agc ccg cgc gta ccg aca gaa gcc				2112
Gly Pro Gly Val Tyr Asp Ile His Ser Pro Arg Val Pro Thr Glu Ala				
	690	695	700	
gaa gtg gag cac ctg ttg cgc aaa gcc atc gag gtt gta ccg gtt gaa				2160
Glu Val Glu His Leu Leu Arg Lys Ala Ile Glu Val Val Pro Val Glu				
705	710	715	720	
cgt ctg tgg gtt aac ccg gac tgc ggc ctg aaa aca cgc ggc tgg aaa				2208
Arg Leu Trp Val Asn Pro Asp Cys Gly Leu Lys Thr Arg Gly Trp Lys				
	725	730	735	
gaa act ctg gaa caa ctc caa gtg atg atg aac gta acc cac aaa ttg				2256
Glu Thr Leu Glu Gln Leu Gln Val Met Met Asn Val Thr His Lys Leu				
	740	745	750	
cgt gcc gaa ttg gcg aaa				2274
Arg Ala Glu Leu Ala Lys				
	755			

<210> 72

<211> 758

<212> PRT

<213> Neisseria meningitidis

<400> 72

Met Thr Thr Leu His Phe Ser Gly Phe Pro Arg Val Gly Ala Phe Arg
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Glu Leu Lys Phe Ala Gln Glu Lys Tyr Trp Arg Lys Glu Ile Ser Glu
20 25 30

Gln Glu Leu Leu Ala Val Ala Lys Asp Leu Arg Glu Lys Asn Trp Lys

35	40	45
His Gln Ala Ala Ala Asn Ala Asp Tyr Val Ala Val Gly Asp Phe Thr		
50	55	60
Phe Tyr Asp His Ile Leu Asp Leu Gln Val Ala Thr Gly Ala Ile Pro		
65	70	75
Ala Arg Phe Gly Phe Asp Ser Gln Asn Leu Ser Leu Glu Gln Phe Phe		
85	90	95
Gln Leu Ala Arg Gly Asn Lys Asp Gln Phe Ala Ile Glu Met Thr Lys		
100	105	110
Trp Phe Asp Thr Asn Tyr His Tyr Leu Val Pro Glu Phe His Ala Asp		
115	120	125
Thr Glu Phe Lys Ala Asn Ala Lys His Tyr Val Gln Gln Leu Gln Glu		
130	135	140
Ala Gln Ala Leu Gly Leu Lys Ala Lys Pro Thr Val Val Gly Pro Leu		
145	150	155
Thr Phe Leu Trp Val Gly Lys Glu Lys Gly Ala Val Glu Phe Asp Arg		
165	170	175
Leu Ser Leu Leu Pro Lys Leu Leu Pro Val Tyr Val Glu Ile Leu Thr		
180	185	190
Ala Leu Val Glu Ala Gly Ala Glu Trp Ile Gln Ile Asp Glu Pro Ala		
195	200	205
Leu Thr Val Asp Leu Pro Lys Glu Trp Val Glu Ala Tyr Lys Asp Val		
210	215	220
Tyr Ala Thr Leu Ser Lys Val Ser Ala Lys Ile Leu Leu Ser Thr Tyr		
225	230	235
Phe Gly Ser Val Ala Glu His Ala Ala Leu Leu Lys Ser Leu Pro Val		
245	250	255
Asp Gly Leu His Ile Asp Leu Val Arg Ala Pro Glu Gln Leu Asp Ala		
260	265	270
Phe Ala Asp Tyr Asp Lys Val Leu Ser Ala Gly Val Ile Asp Gly Arg		
275	280	285
Asn Ile Trp Arg Ala Asn Leu Asn Lys Val Leu Glu Thr Val Glu Leu		

290	295	300
Leu Gln Ala Lys Leu Gly Asp Arg Leu Trp Ile Ser Ser Ser Cys Ser		
305	310	315 320
Leu Leu His Thr Pro Phe Asp Leu Ser Val Glu Glu Lys Leu Lys Ala		
	325	330 335
Asn Lys Pro Asp Leu Tyr Ser Trp Leu Ala Phe Thr Leu Gln Lys Thr		
	340	345 350
Gln Glu Leu Arg Val Leu Lys Ala Ala Leu Asn Glu Gly Arg Asp Ser		
	355	360 365
Val Ala Glu Glu Leu Ala Ala Ser Gln Ala Ala Ala Asp Ser Arg Ala		
	370	375 380
Asn Ser Ser Glu Ile His Arg Ala Asp Val Ala Lys Arg Leu Ala Asp		
385	390	395 400
Leu Pro Ala Asn Ala Asp Gln Arg Lys Ser Pro Phe Ala Asp Arg Ile		
	405	410 415
Lys Ala Gln Gln Ala Trp Leu Asn Leu Pro Leu Leu Pro Thr Thr Asn		
	420	425 430
Ile Gly Ser Phe Pro Gln Thr Thr Glu Ile Arg Gln Ala Arg Ala Ala		
	435	440 445
Phe Lys Lys Gly Glu Leu Ser Ala Ala Asp Tyr Glu Ala Ala Met Lys		
	450	455 460
Lys Glu Ile Ala Leu Val Val Glu Glu Gln Glu Lys Leu Asp Leu Asp		
465	470	475 480
Val Leu Val His Gly Glu Ala Glu Arg Asn Asp Met Val Glu Tyr Phe		
	485	490 495
Gly Glu Leu Leu Ser Gly Phe Ala Phe Thr Gln Tyr Gly Trp Val Gln		
	500	505 510
Ser Tyr Gly Ser Arg Cys Val Lys Pro Pro Ile Ile Phe Gly Asp Val		
	515	520 525
Ser Arg Pro Glu Ala Met Thr Val Ala Trp Ser Thr Tyr Ala Gln Ser		
	530	535 540
Leu Thr Lys Arg Pro Met Lys Gly Met Leu Thr Gly Pro Val Thr Ile		

545 550 555 560
 Leu Gln Trp Ser Phe Val Arg Asn Asp Ile Pro Arg Ser Thr Val Cys
 565 570 575
 Lys Gln Ile Ala Leu Ala Leu Asn Asp Glu Val Leu Asp Leu Glu Lys
 580 585 590
 Ala Gly Ile Lys Val Ile Gln Ile Asp Glu Pro Ala Ile Arg Glu Gly
 595 600 605
 Leu Pro Leu Lys Arg Ala Asp Trp Asp Ala Tyr Leu Asn Trp Ala Gly
 610 615 620
 Glu Ser Phe Arg Leu Ser Ser Thr Gly Cys Glu Asp Ser Thr Gln Ile
 625 630 635 640
 His Thr His Met Cys Tyr Ser Glu Phe Asn Asp Ile Leu Pro Ala Ile
 645 650 655
 Ala Ala Met Asp Ala Asp Val Ile Thr Ile Glu Thr Ser Arg Ser Asp
 660 665 670
 Met Glu Leu Leu Thr Ala Phe Gly Glu Phe Lys Tyr Pro Asn Asp Ile
 675 680 685
 Gly Pro Gly Val Tyr Asp Ile His Ser Pro Arg Val Pro Thr Glu Ala
 690 695 700
 Glu Val Glu His Leu Leu Arg Lys Ala Ile Glu Val Val Pro Val Glu
 705 710 715 720
 Arg Leu Trp Val Asn Pro Asp Cys Gly Leu Lys Thr Arg Gly Trp Lys
 725 730 735
 Glu Thr Leu Glu Gln Leu Gln Val Met Met Asn Val Thr His Lys Leu
 740 745 750
 Arg Ala Glu Leu Ala Lys
 755

<210> 73

<211> 2118

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(2118)

<400> 73

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Met Arg Arg Phe Leu Pro Ile Ala Ala Ile Cys Ala Val Val Leu Leu
  1             5             10             15

tac gga ctg acg gcg gca acc ggc agc acc agt tcg ctg gcg gat tat 96
Tyr Gly Leu Thr Ala Ala Thr Gly Ser Thr Ser Ser Leu Ala Asp Tyr
          20             25             30

ttc tgg tgg att gtt gcg ttc agc gca atg ctg ctg ctg gtg ttg tcc 144
Phe Trp Trp Ile Val Ala Phe Ser Ala Met Leu Leu Leu Val Leu Ser
          35             40             45

gcc gtt ttg gca cgt tat gtc ata ttg ctg ttg aaa gac agg cgc gac 192
Ala Val Leu Ala Arg Tyr Val Ile Leu Leu Leu Lys Asp Arg Arg Asp
          50             55             60

ggc gta ttc ggt tcg cag att gcc aaa cgc ctt tcc ggg atg ttt acg 240
Gly Val Phe Gly Ser Gln Ile Ala Lys Arg Leu Ser Gly Met Phe Thr
          65             70             75             80

ctg gtt gcc gta ctg ccc ggc gtg ttt ctg ttc ggc gtt tcc gca cag 288
Leu Val Ala Val Leu Pro Gly Val Phe Leu Phe Gly Val Ser Ala Gln
          85             90             95

ttt atc aac ggc acg att aat tcg tgg ttc ggc aac gat acc cac gag 336
Phe Ile Asn Gly Thr Ile Asn Ser Trp Phe Gly Asn Asp Thr His Glu
          100             105             110

gcg ctt gaa cgc agc ctc aat ttg agc aag tcc gca ttg aat ctg gcg 384
Ala Leu Glu Arg Ser Leu Asn Leu Ser Lys Ser Ala Leu Asn Leu Ala
          115             120             125

gca gac aac gcc ctt ggc aac gcc atc ccc gtg cag ata gac ctc atc 432
Ala Asp Asn Ala Leu Gly Asn Ala Ile Pro Val Gln Ile Asp Leu Ile
          130             135             140

ggc gcg gct tcc ctg ccc ggg gat atg ggc agg gtg ctg gaa cat tac 480
Gly Ala Ala Ser Leu Pro Gly Asp Met Gly Arg Val Leu Glu His Tyr
          145             150             155             160

gcc ggc agc ggt ttt gcc cag ctt gcc ctg tac aat gcc gca agc ggc 528
Ala Gly Ser Gly Phe Ala Gln Leu Ala Leu Tyr Asn Ala Ala Ser Gly
          165             170             175

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aaa atc gaa aaa agc atc aac ccg cac aag ctc gat cag ccg ttt cca 576
Lys Ile Glu Lys Ser Ile Asn Pro His Lys Leu Asp Gln Pro Phe Pro
      180                      185                      190

ggt aag gcg cgt tgg gaa aaa atc caa cag gcg ggt tcg gtc agg gat 624
Gly Lys Ala Arg Trp Glu Lys Ile Gln Gln Ala Gly Ser Val Arg Asp
      195                      200                      205

ttg gaa agc ata ggc ggc gta ttg tac gcg cag ggc tgg ctg tcg gca 672
Leu Glu Ser Ile Gly Gly Val Leu Tyr Ala Gln Gly Trp Leu Ser Ala
      210                      215                      220

ggt acg cac aac ggg cgc gat tac gcc ttg ttt ttc cgt cag ccg gtt 720
Gly Thr His Asn Gly Arg Asp Tyr Ala Leu Phe Phe Arg Gln Pro Val
225                      230                      235                      240

ccc aaa ggc gtg gca gag gat gcc gtc tta atc gaa aag gca agg gcg 768
Pro Lys Gly Val Ala Glu Asp Ala Val Leu Ile Glu Lys Ala Arg Ala
      245                      250                      255

aaa tat gct gag ttg agt tac agc aaa aaa ggt ttg cag acc ttt ttc 816
Lys Tyr Ala Glu Leu Ser Tyr Ser Lys Lys Gly Leu Gln Thr Phe Phe
      260                      265                      270

ctg gca acc ctg ctg att gcc tcg ctg ctg tcg att ttt ctt gca ctg 864
Leu Ala Thr Leu Leu Ile Ala Ser Leu Leu Ser Ile Phe Leu Ala Leu
      275                      280                      285

gtc atg gca ctg tat ttc gcc cgc cgt ttc gtc gaa ccc gtc cta tcg 912
Val Met Ala Leu Tyr Phe Ala Arg Arg Phe Val Glu Pro Val Leu Ser
      290                      295                      300

ctt gcc gag ggg gcg aag gcg gtg gcg caa ggc gat ttc agc cag acg 960
Leu Ala Glu Gly Ala Lys Ala Val Ala Gln Gly Asp Phe Ser Gln Thr
305                      310                      315                      320

cgc ccc gtg ttg cgc aac gac gag ttc gga cgc ttg acc aag ttg ttc 1008
Arg Pro Val Leu Arg Asn Asp Glu Phe Gly Arg Leu Thr Lys Leu Phe
      325                      330                      335

aac cac atg acc gag cag ctt tcc atc gcc aaa gaa gca gac gag cgc 1056
Asn His Met Thr Glu Gln Leu Ser Ile Ala Lys Glu Ala Asp Glu Arg
      340                      345                      350

aac cgc cgg cgc gag gaa gcc gcc aga cat tat ctc gaa tgc gtg ttg 1104
Asn Arg Arg Arg Glu Glu Ala Ala Arg His Tyr Leu Glu Cys Val Leu
      355                      360                      365

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gag ggg ctg acc acg ggc gtg gtg gtg ttt gac gaa caa ggc tgt ctg	1152
Glu Gly Leu Thr Thr Gly Val Val Val Phe Asp Glu Gln Gly Cys Leu	
370 375 380	
aaa acc ttc aac aaa gcg gcg gaa cag att ttg ggg atg ccg ctt acc	1200
Lys Thr Phe Asn Lys Ala Ala Glu Gln Ile Leu Gly Met Pro Leu Thr	
385 390 395 400	
ccc ctg tgg ggc agc agc cgg cac ggt tgg cac ggc gtt tcg gcg cag	1248
Pro Leu Trp Gly Ser Ser Arg His Gly Trp His Gly Val Ser Ala Gln	
405 410 415	
cag tcc ctg ctt gcc gaa gtg ttt gcc gcc atc ggc gcg gcg gca ggt	1296
Gln Ser Leu Leu Ala Glu Val Phe Ala Ala Ile Gly Ala Ala Ala Gly	
420 425 430	
acg gac aaa ccg gtc cat gtg aaa tat gcc gcg ccg gac gat gcc aaa	1344
Thr Asp Lys Pro Val His Val Lys Tyr Ala Ala Pro Asp Asp Ala Lys	
435 440 445	
atc ctg ctg ggc aag gca acc gtc ctg ccc gaa gac aac ggc aac ggc	1392
Ile Leu Leu Gly Lys Ala Thr Val Leu Pro Glu Asp Asn Gly Asn Gly	
450 455 460	
gtg gta atg gtg att gac gac atc acc gtt ttg ata cac gcg caa aaa	1440
Val Val Met Val Ile Asp Asp Ile Thr Val Leu Ile His Ala Gln Lys	
465 470 475 480	
gaa gcc gcg tgg ggc gaa gtg gca aaa ccg ctg gca cac gaa atc cgc	1488
Glu Ala Ala Trp Gly Glu Val Ala Lys Arg Leu Ala His Glu Ile Arg	
485 490 495	
aat ccg ctc acg ccc atc cag ctt tct gcc gaa ccg ctg gcg tgg aaa	1536
Asn Pro Leu Thr Pro Ile Gln Leu Ser Ala Glu Arg Leu Ala Trp Lys	
500 505 510	
ttg ggc ggg aag ctg gac gag cag gac gcg caa atc ctg aca cgt tcg	1584
Leu Gly Gly Lys Leu Asp Glu Gln Asp Ala Gln Ile Leu Thr Arg Ser	
515 520 525	
acc gac acc atc atc aaa caa gtg gcg gca tta aaa gaa atg gtc gag	1632
Thr Asp Thr Ile Ile Lys Gln Val Ala Ala Leu Lys Glu Met Val Glu	
530 535 540	
gca ttc cgc aat tac gcg cgt tcc cct tcg ctc aaa ttg gaa aat cag	1680
Ala Phe Arg Asn Tyr Ala Arg Ser Pro Ser Leu Lys Leu Glu Asn Gln	
545 550 555 560	

gat ttg aac gcc tta atc ggc gat gtg ttg gca ttg tac gaa gct ggt 1728
 Asp Leu Asn Ala Leu Ile Gly Asp Val Leu Ala Leu Tyr Glu Ala Gly
 565 570 575

ccg tgc cgg ttt gcg gcg gaa ctt gcc ggc gaa ccg ctg atg atg gcg 1776
 Pro Cys Arg Phe Ala Ala Glu Leu Ala Gly Glu Pro Leu Met Met Ala
 580 585 590

gcg gat acg acc gcc atg cgg cag gtg ctg cac aat att ttc aaa aat 1824
 Ala Asp Thr Thr Ala Met Arg Gln Val Leu His Asn Ile Phe Lys Asn
 595 600 605

gcc gcc gaa gcg gcg gaa gaa gcc gat gtg ccc gaa gtc agg gta aaa 1872
 Ala Ala Glu Ala Ala Glu Glu Ala Asp Val Pro Glu Val Arg Val Lys
 610 615 620

tcg gaa gcg ggg cag gac gga cgg att gtc ctg aca gtt tgc gac aac 1920
 Ser Glu Ala Gly Gln Asp Gly Arg Ile Val Leu Thr Val Cys Asp Asn
 625 630 635 640

ggc aag ggg ttc ggc agg gaa atg ctg cac aat gcc ttc gag ccg tat 1968
 Gly Lys Gly Phe Gly Arg Glu Met Leu His Asn Ala Phe Glu Pro Tyr
 645 650 655

gta acg gac aaa ccg gct gga acg gga ttg gga ctg ccc gtg gtg aaa 2016
 Val Thr Asp Lys Pro Ala Gly Thr Gly Leu Gly Leu Pro Val Val Lys
 660 665 670

aaa atc att gaa gaa cac ggc ggc cgc atc agc ctg agc aat cag gat 2064
 Lys Ile Ile Glu Glu His Gly Gly Arg Ile Ser Leu Ser Asn Gln Asp
 675 680 685

gcg ggc ggc gcg tgt gtc aga atc atc ttg cca aaa acg gta gaa act 2112
 Ala Gly Gly Ala Cys Val Arg Ile Ile Leu Pro Lys Thr Val Glu Thr
 690 695 700

tat gcg 2118
 Tyr Ala
 705

<210> 74

<211> 706

<212> PRT

<213> Neisseria meningitidis

<400> 74

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 Tyr Gly Leu Thr Ala Ala Thr Gly Ser Thr Ser Ser Leu Ala Asp Tyr
 20 25 30
 Phe Trp Trp Ile Val Ala Phe Ser Ala Met Leu Leu Leu Val Leu Ser
 35 40 45
 Ala Val Leu Ala Arg Tyr Val Ile Leu Leu Leu Lys Asp Arg Arg Asp
 50 55 60
 Gly Val Phe Gly Ser Gln Ile Ala Lys Arg Leu Ser Gly Met Phe Thr
 65 70 75 80
 Leu Val Ala Val Leu Pro Gly Val Phe Leu Phe Gly Val Ser Ala Gln
 85 90 95
 Phe Ile Asn Gly Thr Ile Asn Ser Trp Phe Gly Asn Asp Thr His Glu
 100 105 110
 Ala Leu Glu Arg Ser Leu Asn Leu Ser Lys Ser Ala Leu Asn Leu Ala
 115 120 125
 Ala Asp Asn Ala Leu Gly Asn Ala Ile Pro Val Gln Ile Asp Leu Ile
 130 135 140
 Gly Ala Ala Ser Leu Pro Gly Asp Met Gly Arg Val Leu Glu His Tyr
 145 150 155 160
 Ala Gly Ser Gly Phe Ala Gln Leu Ala Leu Tyr Asn Ala Ala Ser Gly
 165 170 175
 Lys Ile Glu Lys Ser Ile Asn Pro His Lys Leu Asp Gln Pro Phe Pro
 180 185 190
 Gly Lys Ala Arg Trp Glu Lys Ile Gln Gln Ala Gly Ser Val Arg Asp
 195 200 205
 Leu Glu Ser Ile Gly Gly Val Leu Tyr Ala Gln Gly Trp Leu Ser Ala
 210 215 220
 Gly Thr His Asn Gly Arg Asp Tyr Ala Leu Phe Phe Arg Gln Pro Val
 225 230 235 240
 Pro Lys Gly Val Ala Glu Asp Ala Val Leu Ile Glu Lys Ala Arg Ala
 245 250 255

Lys Tyr Ala Glu Leu Ser Tyr Ser Lys Lys Gly Leu Gln Thr Phe Phe
 260 265 270

Leu Ala Thr Leu Leu Ile Ala Ser Leu Leu Ser Ile Phe Leu Ala Leu
 275 280 285

Val Met Ala Leu Tyr Phe Ala Arg Arg Phe Val Glu Pro Val Leu Ser
 290 295 300

Leu Ala Glu Gly Ala Lys Ala Val Ala Gln Gly Asp Phe Ser Gln Thr
 305 310 315 320

Arg Pro Val Leu Arg Asn Asp Glu Phe Gly Arg Leu Thr Lys Leu Phe
 325 330 335

Asn His Met Thr Glu Gln Leu Ser Ile Ala Lys Glu Ala Asp Glu Arg
 340 345 350

Asn Arg Arg Arg Glu Glu Ala Ala Arg His Tyr Leu Glu Cys Val Leu
 355 360 365

Glu Gly Leu Thr Thr Gly Val Val Val Phe Asp Glu Gln Gly Cys Leu
 370 375 380

Lys Thr Phe Asn Lys Ala Ala Glu Gln Ile Leu Gly Met Pro Leu Thr
 385 390 395 400

Pro Leu Trp Gly Ser Ser Arg His Gly Trp His Gly Val Ser Ala Gln
 405 410 415

Gln Ser Leu Leu Ala Glu Val Phe Ala Ala Ile Gly Ala Ala Ala Gly
 420 425 430

Thr Asp Lys Pro Val His Val Lys Tyr Ala Ala Pro Asp Asp Ala Lys
 435 440 445

Ile Leu Leu Gly Lys Ala Thr Val Leu Pro Glu Asp Asn Gly Asn Gly
 450 455 460

Val Val Met Val Ile Asp Asp Ile Thr Val Leu Ile His Ala Gln Lys
 465 470 475 480

Glu Ala Ala Trp Gly Glu Val Ala Lys Arg Leu Ala His Glu Ile Arg
 485 490 495

Asn Pro Leu Thr Pro Ile Gln Leu Ser Ala Glu Arg Leu Ala Trp Lys
 500 505 510

Leu Gly Gly Lys Leu Asp Glu Gln Asp Ala Gln Ile Leu Thr Arg Ser
 515 520 525

Thr Asp Thr Ile Ile Lys Gln Val Ala Ala Leu Lys Glu Met Val Glu
 530 535 540

Ala Phe Arg Asn Tyr Ala Arg Ser Pro Ser Leu Lys Leu Glu Asn Gln
 545 550 555 560

Asp Leu Asn Ala Leu Ile Gly Asp Val Leu Ala Leu Tyr Glu Ala Gly
 565 570 575

Pro Cys Arg Phe Ala Ala Glu Leu Ala Gly Glu Pro Leu Met Met Ala
 580 585 590

Ala Asp Thr Thr Ala Met Arg Gln Val Leu His Asn Ile Phe Lys Asn
 595 600 605

Ala Ala Glu Ala Ala Glu Glu Ala Asp Val Pro Glu Val Arg Val Lys
 610 615 620

Ser Glu Ala Gly Gln Asp Gly Arg Ile Val Leu Thr Val Cys Asp Asn
 625 630 635 640

Gly Lys Gly Phe Gly Arg Glu Met Leu His Asn Ala Phe Glu Pro Tyr
 645 650 655

Val Thr Asp Lys Pro Ala Gly Thr Gly Leu Gly Leu Pro Val Val Lys
 660 665 670

Lys Ile Ile Glu Glu His Gly Gly Arg Ile Ser Leu Ser Asn Gln Asp
 675 680 685

Ala Gly Gly Ala Cys Val Arg Ile Ile Leu Pro Lys Thr Val Glu Thr
 690 695 700

Tyr Ala
 705

<210> 75

<211> 1686

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1686)

<400> 75

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Met Leu Gln Ser Glu Asn Ser Lys Asn Leu Ile Ser Cys Ser Ile Lys
  1             5             10             15

gct tta cca atc atg att tct acc aac ggc atc acc atg cag ttc ggc 96
Ala Leu Pro Ile Met Ile Ser Thr Asn Gly Ile Thr Met Gln Phe Gly
      20             25             30

gcg aag ccg ctg ttt gaa aac gta tcc gtt aaa ttc ggc gaa ggc aac 144
Ala Lys Pro Leu Phe Glu Asn Val Ser Val Lys Phe Gly Glu Gly Asn
      35             40             45

cgc tac ggt ttg atc ggc gcg aac ggc tca ggc aaa tcc acc ttc atg 192
Arg Tyr Gly Leu Ile Gly Ala Asn Gly Ser Gly Lys Ser Thr Phe Met
      50             55             60

aaa atc ctc ggc ggc gat ttg gaa cag aca gcc ggc gaa gtg gcg att 240
Lys Ile Leu Gly Gly Asp Leu Glu Gln Thr Ala Gly Glu Val Ala Ile
      65             70             75             80

gaa aac ggc gtg cgt ttg ggt aaa ttg cgc caa gac cag ttt gcc tac 288
Glu Asn Gly Val Arg Leu Gly Lys Leu Arg Gln Asp Gln Phe Ala Tyr
      85             90             95

gaa gat atg cgc gtg ctg gac gtg gta atg atg ggt cac acc gaa atg 336
Glu Asp Met Arg Val Leu Asp Val Val Met Met Gly His Thr Glu Met
      100             105             110

tgg gcg gcg atg acc gaa cgt gat gcg att tac gcc aat ccc gaa gcc 384
Trp Ala Ala Met Thr Glu Arg Asp Ala Ile Tyr Ala Asn Pro Glu Ala
      115             120             125

acc gaa gac gac tac atg aaa gcc gca gaa ctg gaa gcc aag ttc gcc 432
Thr Glu Asp Asp Tyr Met Lys Ala Ala Glu Leu Glu Ala Lys Phe Ala
      130             135             140

gaa tac gac ggc tac acc gcc gaa gcg cgt gcc gcc gag ttg ttg agc 480
Glu Tyr Asp Gly Tyr Thr Ala Glu Ala Arg Ala Ala Glu Leu Leu Ser
      145             150             155             160

ggc gtg ggc att tcc gaa gat ttg cac aat gca acc atg gca gaa gtt 528
Gly Val Gly Ile Ser Glu Asp Leu His Asn Ala Thr Met Ala Glu Val
      165             170             175

gcc ccg ggc ttc aaa ctg cgc gta ttg ctg gcg caa gcc ctg ttc tcc 576

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Ala Pro Gly Phe Lys Leu Arg Val Leu Leu Ala Gln Ala Leu Phe Ser	
180 185 190	
aag ccg gat gta ttg ctc ttg gac gaa ccg acc aat aac ttg gac att	624
Lys Pro Asp Val Leu Leu Leu Asp Glu Pro Thr Asn Asn Leu Asp Ile	
195 200 205	
aat acc atc cgt tgg ttg gaa ggc gtg ttg aac cag tac gac tcc acg	672
Asn Thr Ile Arg Trp Leu Glu Gly Val Leu Asn Gln Tyr Asp Ser Thr	
210 215 220	
atg att atc atc tcg cac gac cgc cac ttc ttg aac gaa gtc tgc acg	720
Met Ile Ile Ile Ser His Asp Arg His Phe Leu Asn Glu Val Cys Thr	
225 230 235 240	
cat atg gca gat ttg gat tac aac acc atc acc atc tat ccg ggc aac	768
His Met Ala Asp Leu Asp Tyr Asn Thr Ile Thr Ile Tyr Pro Gly Asn	
245 250 255	
tac gac gac tat atg ctc gcc tcc gcc caa tcg cgc gaa cgc gcc ctg	816
Tyr Asp Asp Tyr Met Leu Ala Ser Ala Gln Ser Arg Glu Arg Ala Leu	
260 265 270	
aaa gac aat gcc aag gcg aag gaa aaa ttg cag gaa ttg caa gaa ttt	864
Lys Asp Asn Ala Lys Ala Lys Glu Lys Leu Gln Glu Leu Gln Glu Phe	
275 280 285	
gtg gcg cgg ttc agt gcc aac aaa tcc aaa gcc cgt cag gca acc agc	912
Val Ala Arg Phe Ser Ala Asn Lys Ser Lys Ala Arg Gln Ala Thr Ser	
290 295 300	
cgt ctg aaa cag gca gac aaa atc aaa tcg gag atg gtc gag gtc aaa	960
Arg Leu Lys Gln Ala Asp Lys Ile Lys Ser Glu Met Val Glu Val Lys	
305 310 315 320	
ccc tct acc cgt caa aac ccg tat atc cgt ttt gaa gcc gat gaa aaa	1008
Pro Ser Thr Arg Gln Asn Pro Tyr Ile Arg Phe Glu Ala Asp Glu Lys	
325 330 335	
gcc aag ctg cac cgt cag gct gtg gaa gtt gaa aaa ctg gcg aaa cgc	1056
Ala Lys Leu His Arg Gln Ala Val Glu Val Glu Lys Leu Ala Lys Arg	
340 345 350	
ttt gaa acc cag ttg ttt aaa aac ctg aac ttt atc ctt gaa gcg gga	1104
Phe Glu Thr Gln Leu Phe Lys Asn Leu Asn Phe Ile Leu Glu Ala Gly	
355 360 365	
caa cgc ctc gcc atc atc ggc ccg aac ggc gcg ggc aaa tcc acc ctg	1152

Gln Arg Leu Ala Ile Ile Gly Pro Asn Gly Ala Gly Lys Ser Thr Leu
 370 375 380

 ctg aaa ctc ttg gcc ggc gcg tac aac ccc gaa tat tca gac ggt ctg 1200
 Leu Lys Leu Leu Ala Gly Ala Tyr Asn Pro Glu Tyr Ser Asp Gly Leu
 385 390 395 400

 ctg ccg gac gaa ggc agc atc aaa tgg gcg gaa aaa gcc agc gtc ggc 1248
 Leu Pro Asp Glu Gly Ser Ile Lys Trp Ala Glu Lys Ala Ser Val Gly
 405 410 415

 tac tat ccg caa gat cac gaa aac gac ttc gat gtc gat atg gat ttg 1296
 Tyr Tyr Pro Gln Asp His Glu Asn Asp Phe Asp Val Asp Met Asp Leu
 420 425 430

 agc gaa tgg atg cgc caa tgg ggg cag gac ggc gac gac gaa caa gtc 1344
 Ser Glu Trp Met Arg Gln Trp Gly Gln Asp Gly Asp Asp Glu Gln Val
 435 440 445

 atc cgc ggc act ttg ggg cgt ttg ctc ttc ggc agt aac gat gtc gtg 1392
 Ile Arg Gly Thr Leu Gly Arg Leu Leu Phe Gly Ser Asn Asp Val Val
 450 455 460

 aaa aaa gtg aag gtt ctc tcc ggt ggt gaa aaa ggc cgt atg ctt tac 1440
 Lys Lys Val Lys Val Leu Ser Gly Gly Glu Lys Gly Arg Met Leu Tyr
 465 470 475 480

 ggc aaa ctg ttg ctg ttg aaa ccc aat gtc tta gtc atg gat gaa ccg 1488
 Gly Lys Leu Leu Leu Lys Pro Asn Val Leu Val Met Asp Glu Pro
 485 490 495

 acc aac cac atg gac atg gaa agc atc gaa tcc ttg aac atg gcg ctg 1536
 Thr Asn His Met Asp Met Glu Ser Ile Glu Ser Leu Asn Met Ala Leu
 500 505 510

 gaa aaa tac aac ggc acg ctg att ttc gtc tcc cac gac cgt cag ttc 1584
 Glu Lys Tyr Asn Gly Thr Leu Ile Phe Val Ser His Asp Arg Gln Phe
 515 520 525

 gtt tct tcc ctg gct acc caa atc att gaa ttg gac ggc aaa ggc gga 1632
 Val Ser Ser Leu Ala Thr Gln Ile Ile Glu Leu Asp Gly Lys Gly Gly
 530 535 540

 tat gaa cac tac ttg ggc gat tac gaa agc tat ctc gag aaa aaa ggc 1680
 Tyr Glu His Tyr Leu Gly Asp Tyr Glu Ser Tyr Leu Glu Lys Lys Gly
 545 550 555 560

 gta gca 1686

Val Ala

<210> 76

<211> 562

<212> PRT

<213> Neisseria meningitidis

<400> 76

Met Leu Gln Ser Glu Asn Ser Lys Asn Leu Ile Ser Cys Ser Ile Lys

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Ala Leu Pro Ile Met Ile Ser Thr Asn Gly Ile Thr Met Gln Phe Gly

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Ala Lys Pro Leu Phe Glu Asn Val Ser Val Lys Phe Gly Glu Gly Asn

35 40 45

Arg Tyr Gly Leu Ile Gly Ala Asn Gly Ser Gly Lys Ser Thr Phe Met

50 55 60

Lys Ile Leu Gly Gly Asp Leu Glu Gln Thr Ala Gly Glu Val Ala Ile

65 70 75 80

Glu Asn Gly Val Arg Leu Gly Lys Leu Arg Gln Asp Gln Phe Ala Tyr

85 90 95

Glu Asp Met Arg Val Leu Asp Val Val Met Met Gly His Thr Glu Met

100 105 110

Trp Ala Ala Met Thr Glu Arg Asp Ala Ile Tyr Ala Asn Pro Glu Ala

115 120 125

Thr Glu Asp Asp Tyr Met Lys Ala Ala Glu Leu Glu Ala Lys Phe Ala

130 135 140

Glu Tyr Asp Gly Tyr Thr Ala Glu Ala Arg Ala Ala Glu Leu Leu Ser

145 150 155 160

Gly Val Gly Ile Ser Glu Asp Leu His Asn Ala Thr Met Ala Glu Val

165 170 175

Ala Pro Gly Phe Lys Leu Arg Val Leu Leu Ala Gln Ala Leu Phe Ser

180 185 190

Lys Pro Asp Val Leu Leu Leu Asp Glu Pro Thr Asn Asn Leu Asp Ile

195 200 205

Asn Thr Ile Arg Trp Leu Glu Gly Val Leu Asn Gln Tyr Asp Ser Thr
 210 215 220

Met Ile Ile Ile Ser His Asp Arg His Phe Leu Asn Glu Val Cys Thr
 225 230 235 240

His Met Ala Asp Leu Asp Tyr Asn Thr Ile Thr Ile Tyr Pro Gly Asn
 245 250 255

Tyr Asp Asp Tyr Met Leu Ala Ser Ala Gln Ser Arg Glu Arg Ala Leu
 260 265 270

Lys Asp Asn Ala Lys Ala Lys Glu Lys Leu Gln Glu Leu Gln Glu Phe
 275 280 285

Val Ala Arg Phe Ser Ala Asn Lys Ser Lys Ala Arg Gln Ala Thr Ser
 290 295 300

Arg Leu Lys Gln Ala Asp Lys Ile Lys Ser Glu Met Val Glu Val Lys
 305 310 315 320

Pro Ser Thr Arg Gln Asn Pro Tyr Ile Arg Phe Glu Ala Asp Glu Lys
 325 330 335

Ala Lys Leu His Arg Gln Ala Val Glu Val Glu Lys Leu Ala Lys Arg
 340 345 350

Phe Glu Thr Gln Leu Phe Lys Asn Leu Asn Phe Ile Leu Glu Ala Gly
 355 360 365

Gln Arg Leu Ala Ile Ile Gly Pro Asn Gly Ala Gly Lys Ser Thr Leu
 370 375 380

Leu Lys Leu Leu Ala Gly Ala Tyr Asn Pro Glu Tyr Ser Asp Gly Leu
 385 390 395 400

Leu Pro Asp Glu Gly Ser Ile Lys Trp Ala Glu Lys Ala Ser Val Gly
 405 410 415

Tyr Tyr Pro Gln Asp His Glu Asn Asp Phe Asp Val Asp Met Asp Leu
 420 425 430

Ser Glu Trp Met Arg Gln Trp Gly Gln Asp Gly Asp Asp Glu Gln Val
 435 440 445

Ile Arg Gly Thr Leu Gly Arg Leu Leu Phe Gly Ser Asn Asp Val Val
 450 455 460

Lys Lys Val Lys Val Leu Ser Gly Gly Glu Lys Gly Arg Met Leu Tyr
 465 470 475 480

Gly Lys Leu Leu Leu Lys Pro Asn Val Leu Val Met Asp Glu Pro
 485 490 495

Thr Asn His Met Asp Met Glu Ser Ile Glu Ser Leu Asn Met Ala Leu
 500 505 510

Glu Lys Tyr Asn Gly Thr Leu Ile Phe Val Ser His Asp Arg Gln Phe
 515 520 525

Val Ser Ser Leu Ala Thr Gln Ile Ile Glu Leu Asp Gly Lys Gly Gly
 530 535 540

Tyr Glu His Tyr Leu Gly Asp Tyr Glu Ser Tyr Leu Glu Lys Lys Gly
 545 550 555 560

Val Ala

<210> 77

<211> 1773

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1773)

<400> 77

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ggt tgc gcc cac ctg att gcg cgc ggt acg gag gaa gtg ccg cag tat 96
 Gly Cys Ala His Leu Ile Ala Arg Gly Thr Glu Glu Val Pro Gln Tyr
 20 25 30

gat gtt gcg gag gcg gac acc gat gcc gaa gcc gaa cgt ttc gat gcc 144
 Asp Val Ala Glu Ala Asp Thr Asp Ala Glu Ala Glu Arg Phe Asp Ala
 35 40 45

gcc gtc aaa gcc acg cgc aaa gag ttg gaa cag ctc cgc agc gcg att 192
 Ala Val Lys Ala Thr Arg Lys Glu Leu Glu Gln Leu Arg Ser Ala Ile
 50 55 60

151

gat ttg aac aag ctc aaa aaa acc gcc gcc gcc acc gct gac ggg gtc	816
Asp Leu Asn Lys Leu Lys Lys Thr Ala Ala Ala Thr Ala Asp Gly Val	
260 265 270	
tgc atc gag ctt gtg ggc aat ata gaa tcc gcc gaa gac gtg aaa ccg	864
Cys Ile Glu Leu Val Gly Asn Ile Glu Ser Ala Glu Asp Val Lys Pro	
275 280 285	
ctg cac aac ctc ggc gca gac ggc atc ggg ctg ttc cgc agc gag ttt	912
Leu His Asn Leu Gly Ala Asp Gly Ile Gly Leu Phe Arg Ser Glu Phe	
290 295 300	
ctt tac ctg aac cgc gat acg atg ccg tct gaa gac gag cag tac gaa	960
Leu Tyr Leu Asn Arg Asp Thr Met Pro Ser Glu Asp Glu Gln Tyr Glu	
305 310 315 320	
gtg tac agc gcg att gtc aaa aaa atg aaa ggc aaa agc gta acg ata	1008
Val Tyr Ser Ala Ile Val Lys Lys Met Lys Gly Lys Ser Val Thr Ile	
325 330 335	
cgg aca gtc gat ttg ggt gtg gac aaa aac ccg cgc tgg ttc ggg aaa	1056
Arg Thr Val Asp Leu Gly Val Asp Lys Asn Pro Arg Trp Phe Gly Lys	
340 345 350	
aac agc acg ccc aac ggc agc ctc aac ccc gcg ctg ggc atg acc ggc	1104
Asn Ser Thr Pro Asn Gly Ser Leu Asn Pro Ala Leu Gly Met Thr Gly	
355 360 365	
atc cgc ctg tgc ctt gcc gaa ccg gtc atg ttc cgc acc cag atg cgc	1152
Ile Arg Leu Cys Leu Ala Glu Pro Val Met Phe Arg Thr Gln Met Arg	
370 375 380	
gcc atc ctc cgc gcc gcc gcc cac ggc ccc gtg cgg atg atg tgg ccg	1200
Ala Ile Leu Arg Ala Ala Ala His Gly Pro Val Arg Met Met Trp Pro	
385 390 395 400	
atg att acc tcc gta tcc gaa gtg cgc cag tgc ctc atc cac ctc gac	1248
Met Ile Thr Ser Val Ser Glu Val Arg Gln Cys Leu Ile His Leu Asp	
405 410 415	
acc gcg caa cgc cag ctt gcc gaa cgc ggc gat gcc ttc ggt aaa gtc	1296
Thr Ala Gln Arg Gln Leu Ala Glu Arg Gly Asp Ala Phe Gly Lys Val	
420 425 430	
ggc atc ggc tgt atg att gaa att ccg tct gcc gcg ctg acc gtc ggc	1344
Gly Ile Gly Cys Met Ile Glu Ile Pro Ser Ala Ala Leu Thr Val Gly	
435 440 445	

agt att ttg aaa ctg gtc gat ttc atc tcc gtc ggt acc aac gac ctg 1392
 Ser Ile Leu Lys Leu Val Asp Phe Ile Ser Val Gly Thr Asn Asp Leu
 450 455 460

att caa tac atc ttg tcc gtc gat cgc ggc gac gac agc gtc agc cac 1440
 Ile Gln Tyr Ile Leu Ser Val Asp Arg Gly Asp Asp Ser Val Ser His
 465 470 475 480

ctc tac cag ccc ggc cat ccc gcc gtg ctg aaa atg ctg caa cac gtc 1488
 Leu Tyr Gln Pro Gly His Pro Ala Val Leu Lys Met Leu Gln His Val
 485 490 495

atc cgt acc gcc aac cgc atg gac aaa gac gta tcc gta tgc ggc gag 1536
 Ile Arg Thr Ala Asn Arg Met Asp Lys Asp Val Ser Val Cys Gly Glu
 500 505 510

atg gcg ggc gat acc gcg ttt acc cgc gtt tta ttg ggt atg ggg ctg 1584
 Met Ala Gly Asp Thr Ala Phe Thr Arg Val Leu Leu Gly Met Gly Leu
 515 520 525

cgc cgt ttt tcc atg aac ccc aac aac atc ctg ccc gtc aaa aac atc 1632
 Arg Arg Phe Ser Met Asn Pro Asn Asn Ile Leu Pro Val Lys Asn Ile
 530 535 540

att ctg cac agc aat gtc gga cag ctc gaa agc gat att gtg aaa gtc 1680
 Ile Leu His Ser Asn Val Gly Gln Leu Glu Ser Asp Ile Val Lys Val
 545 550 555 560

atc cgc tgc gaa gac gaa gag aag tcg gaa aag ctg atc aaa cag atg 1728
 Ile Arg Cys Glu Asp Glu Glu Lys Ser Glu Lys Leu Ile Lys Gln Met
 565 570 575

aac agc gtg tct gtc gag gaa gaa gcc gac ttc aag ggg cgg aaa 1773
 Asn Ser Val Ser Val Glu Glu Glu Ala Asp Phe Lys Gly Arg Lys
 580 585 590

<210> 78

<211> 591

<212> PRT

<213> Neisseria meningitidis

<400> 78

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35	40	45
Ala Val Lys Ala Thr Arg Lys Glu Leu Glu Gln Leu Arg Ser Ala Ile		
50	55	60
Pro Glu Asn Ala Pro Thr Glu Leu Gly Ala Phe Ile Ser Leu His Leu		
65	70	75
Met Leu Leu Thr Asp Val Thr Leu Ser Arg Glu Pro Val Asp Ile Leu		
85	90	95
Arg Glu Gln Lys Ile Asn Ala Glu Trp Ala Leu Lys Gln Gln Ser Asp		
100	105	110
Lys Leu Ala Ala Gln Phe Asp Asn Met Asp Asp Ala Tyr Leu Arg Glu		
115	120	125
Arg Lys Gln Asp Met Leu Gln Val Val Arg Arg Ile His Asn Asn Leu		
130	135	140
Ile Gly Gln Gly Asn Glu Leu Glu Val Ala Asp Asn Leu Phe Asp Glu		
145	150	155
Thr Val Leu Ile Ala Asn Asp Leu Ser Pro Ala Asp Thr Val Leu Phe		
165	170	175
Lys Glu Gln Arg Ile Ala Ala Phe Val Thr Asp Ala Gly Gly Pro Thr		
180	185	190
Gly His Thr Ala Ile Leu Gly Arg Ser Leu Asp Ile Pro Ser Val Val		
195	200	205
Gly Leu His Asn Ala Arg Lys Leu Ile Thr Glu Gly Glu Thr Val Ile		
210	215	220
Val Asp Gly Ile Asn Gly Val Leu Ile Ile Ala Pro Asp Glu Ser Val		
225	230	235
Leu Asn Glu Tyr Arg Arg Arg Ala Arg Glu Tyr Arg Ser His Lys Arg		
245	250	255
Asp Leu Asn Lys Leu Lys Lys Thr Ala Ala Ala Thr Ala Asp Gly Val		
260	265	270
Cys Ile Glu Leu Val Gly Asn Ile Glu Ser Ala Glu Asp Val Lys Pro		

275	280	285
Leu His Asn Leu Gly Ala Asp Gly Ile Gly Leu Phe Arg Ser Glu Phe		
290	295	300
Leu Tyr Leu Asn Arg Asp Thr Met Pro Ser Glu Asp Glu Gln Tyr Glu		
305	310	315
Val Tyr Ser Ala Ile Val Lys Lys Met Lys Gly Lys Ser Val Thr Ile		
	325	330
Arg Thr Val Asp Leu Gly Val Asp Lys Asn Pro Arg Trp Phe Gly Lys		
	340	345
Asn Ser Thr Pro Asn Gly Ser Leu Asn Pro Ala Leu Gly Met Thr Gly		
	355	360
Ile Arg Leu Cys Leu Ala Glu Pro Val Met Phe Arg Thr Gln Met Arg		
	370	375
Ala Ile Leu Arg Ala Ala Ala His Gly Pro Val Arg Met Met Trp Pro		
385	390	395
Met Ile Thr Ser Val Ser Glu Val Arg Gln Cys Leu Ile His Leu Asp		
	405	410
Thr Ala Gln Arg Gln Leu Ala Glu Arg Gly Asp Ala Phe Gly Lys Val		
	420	425
Gly Ile Gly Cys Met Ile Glu Ile Pro Ser Ala Ala Leu Thr Val Gly		
	435	440
Ser Ile Leu Lys Leu Val Asp Phe Ile Ser Val Gly Thr Asn Asp Leu		
	450	455
Ile Gln Tyr Ile Leu Ser Val Asp Arg Gly Asp Asp Ser Val Ser His		
465	470	475
Leu Tyr Gln Pro Gly His Pro Ala Val Leu Lys Met Leu Gln His Val		
	485	490
Ile Arg Thr Ala Asn Arg Met Asp Lys Asp Val Ser Val Cys Gly Glu		
	500	505
Met Ala Gly Asp Thr Ala Phe Thr Arg Val Leu Leu Gly Met Gly Leu		
	515	520
Arg Arg Phe Ser Met Asn Pro Asn Asn Ile Leu Pro Val Lys Asn Ile		

530 535 540
 Ile Leu His Ser Asn Val Gly Gln Leu Glu Ser Asp Ile Val Lys Val
 545 550 555 560
 Ile Arg Cys Glu Asp Glu Glu Lys Ser Glu Lys Leu Ile Lys Gln Met
 565 570 575
 Asn Ser Val Ser Val Glu Glu Glu Ala Asp Phe Lys Gly Arg Lys
 580 585 590

<210> 79
 <211> 1062
 <212> DNA
 <213> Neisseria meningitidis

<220>
 <221> CDS
 <222> (1)..(1062)

<400> 79
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 1 5 10 15
 ggt gga ggc gaa caa tat gtc tat gat gtt tca aaa gca ttg ggg ctt 96
 Gly Gly Gly Glu Gln Tyr Val Tyr Asp Val Ser Lys Ala Leu Gly Leu
 20 25 30
 cgg gac tgc aca atg ttt acc gcc gtc aat aaa aat aat gaa ttg atg 144
 Arg Asp Cys Thr Met Phe Thr Ala Val Asn Lys Asn Asn Glu Leu Met
 35 40 45
 cac agg cga ttt tcc gaa gtt tct tcc gtt ttc acg aca cgc ctt cac 192
 His Arg Arg Phe Ser Glu Val Ser Ser Val Phe Thr Thr Arg Leu His
 50 55 60
 acg ctc aac ggg ctg ttt tcg ctc tac gca ctt acc cgc ttt atc cgg 240
 Thr Leu Asn Gly Leu Phe Ser Leu Tyr Ala Leu Thr Arg Phe Ile Arg
 65 70 75 80
 aaa aac cgc att tcc cac ctg atg ata cac acc ggc aaa att gcc gcc 288
 Lys Asn Arg Ile Ser His Leu Met Ile His Thr Gly Lys Ile Ala Ala
 85 90 95
 tta tcc ata ctt ttg aaa aaa ctg acc ggg gtg cgc ctg ata ttt gtc 336

Leu Ser Ile	Leu Leu Lys Lys	Leu Thr Gly Val	Arg Leu Ile Phe Val	
100		105	110	
aaa cat aat gtt gtc gcc aac	aaa act gat ttt tat cac cgc ctg ata	384		
Lys His Asn Val Val Ala Asn	Lys Thr Asp Phe Tyr His Arg Leu Ile			
115	120	125		
cag aaa aac acc gac cgc ttt att tgc gtt tcc cgt ctg gtt tac gat	432			
Gln Lys Asn Thr Asp Arg Phe Ile Cys Val Ser Arg Leu Val Tyr Asp				
130	135	140		
gtg caa acc gcc gac aat ccc ttt aaa gaa aaa tac cgg att gtt cat	480			
Val Gln Thr Ala Asp Asn Pro Phe Lys Glu Lys Tyr Arg Ile Val His				
145	150	155	160	
aac ggt atc gat acc ggc cgt ttc cct ccc cct caa gaa aaa ccc gac	528			
Asn Gly Ile Asp Thr Gly Arg Phe Pro Pro Pro Gln Glu Lys Pro Asp				
165	170	175		
agc cgt ttt ttt acc gtc gcc tac gcc ggc agg atc agt ccg gaa aaa	576			
Ser Arg Phe Phe Thr Val Ala Tyr Ala Gly Arg Ile Ser Pro Glu Lys				
180	185	190		
gga ttg gaa aac ctg att gaa gcc tgt gtg ata ctg cat cgg aaa tat	624			
Gly Leu Glu Asn Leu Ile Glu Ala Cys Val Ile Leu His Arg Lys Tyr				
195	200	205		
cct caa atc agg ctc aaa ttg gca ggc gac gga cat ccg gat tat atg	672			
Pro Gln Ile Arg Leu Lys Leu Ala Gly Asp Gly His Pro Asp Tyr Met				
210	215	220		
tgt cgc ctg aag cgg gac gta tct gct tcg gga gca gaa cca ttt gtt	720			
Cys Arg Leu Lys Arg Asp Val Ser Ala Ser Gly Ala Glu Pro Phe Val				
225	230	235	240	
tct ttt gaa ggc ttt acc gaa aaa ctt gct tcg ttt tac cgc caa agc	768			
Ser Phe Glu Gly Phe Thr Glu Lys Leu Ala Ser Phe Tyr Arg Gln Ser				
245	250	255		
gat gtc gtg gtt ttg ccc agc ctc gtc ccg gag gca ttc ggt ttg tca	816			
Asp Val Val Val Leu Pro Ser Leu Val Pro Glu Ala Phe Gly Leu Ser				
260	265	270		
tta tgc gag gcg atg tac tgc cga acg gcg gtg att tcc aat act ttg	864			
Leu Cys Glu Ala Met Tyr Cys Arg Thr Ala Val Ile Ser Asn Thr Leu				
275	280	285		
ggc gcg caa aag gaa att atc gaa cat cat caa tcg ggc att ctg ctg	912			

Gly Ala Gln Lys Glu Ile Ile Glu His His Gln Ser Gly Ile Leu Leu
 290 295 300

gat agg ctg aca cct gaa tct ttg gcg gac gaa atc gaa cgc ctc gtc 960
 Asp Arg Leu Thr Pro Glu Ser Leu Ala Asp Glu Ile Glu Arg Leu Val
 305 310 315 320

tta aac cct gaa acg aaa aac gca ctg gca acg gca gga cat caa tgc 1008
 Leu Asn Pro Glu Thr Lys Asn Ala Leu Ala Thr Ala Gly His Gln Cys
 325 330 335

gtc gcc aac cgt ttt acc atc aac cat acc gcc gac aaa tta ttg gat 1056
 Val Ala Asn Arg Phe Thr Ile Asn His Thr Ala Asp Lys Leu Leu Asp
 340 345 350

gca ata 1062
 Ala Ile

<210> 80

<211> 354

<212> PRT

<213> Neisseria meningitidis

<400> 80

Met Glu Lys Glu Phe Arg Ile Leu Asn Ile Val Ser Ala Lys Ile Trp
 1 5 10 15

Gly Gly Gly Glu Gln Tyr Val Tyr Asp Val Ser Lys Ala Leu Gly Leu
 20 25 30

Arg Asp Cys Thr Met Phe Thr Ala Val Asn Lys Asn Asn Glu Leu Met
 35 40 45

His Arg Arg Phe Ser Glu Val Ser Ser Val Phe Thr Thr Arg Leu His
 50 55 60

Thr Leu Asn Gly Leu Phe Ser Leu Tyr Ala Leu Thr Arg Phe Ile Arg
 65 70 75 80

Lys Asn Arg Ile Ser His Leu Met Ile His Thr Gly Lys Ile Ala Ala
 85 90 95

Leu Ser Ile Leu Leu Lys Lys Leu Thr Gly Val Arg Leu Ile Phe Val
 100 105 110

Lys His Asn Val Val Ala Asn Lys Thr Asp Phe Tyr His Arg Leu Ile
 115 120 125

Gln Lys Asn Thr Asp Arg Phe Ile Cys Val Ser Arg Leu Val Tyr Asp
 130 135 140
 Val Gln Thr Ala Asp Asn Pro Phe Lys Glu Lys Tyr Arg Ile Val His
 145 150 155 160
 Asn Gly Ile Asp Thr Gly Arg Phe Pro Pro Pro Gln Glu Lys Pro Asp
 165 170 175
 Ser Arg Phe Phe Thr Val Ala Tyr Ala Gly Arg Ile Ser Pro Glu Lys
 180 185 190
 Gly Leu Glu Asn Leu Ile Glu Ala Cys Val Ile Leu His Arg Lys Tyr
 195 200 205
 Pro Gln Ile Arg Leu Lys Leu Ala Gly Asp Gly His Pro Asp Tyr Met
 210 215 220
 Cys Arg Leu Lys Arg Asp Val Ser Ala Ser Gly Ala Glu Pro Phe Val
 225 230 235 240
 Ser Phe Glu Gly Phe Thr Glu Lys Leu Ala Ser Phe Tyr Arg Gln Ser
 245 250 255
 Asp Val Val Val Leu Pro Ser Leu Val Pro Glu Ala Phe Gly Leu Ser
 260 265 270
 Leu Cys Glu Ala Met Tyr Cys Arg Thr Ala Val Ile Ser Asn Thr Leu
 275 280 285
 Gly Ala Gln Lys Glu Ile Ile Glu His His Gln Ser Gly Ile Leu Leu
 290 295 300
 Asp Arg Leu Thr Pro Glu Ser Leu Ala Asp Glu Ile Glu Arg Leu Val
 305 310 315 320
 Leu Asn Pro Glu Thr Lys Asn Ala Leu Ala Thr Ala Gly His Gln Cys
 325 330 335
 Val Ala Asn Arg Phe Thr Ile Asn His Thr Ala Asp Lys Leu Leu Asp
 340 345 350
 Ala Ile

<210> 81

<211> 795

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(795)

<400> 81

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atg gcg gct gtt ttt tct gta tgt gcc tgt ttt atg tgt tcc tgc ctt 48
Met Ala Ala Val Phe Ser Val Cys Ala Cys Phe Met Cys Ser Cys Leu
  1           5           10          15

gtt gtc aaa aat acc gtt atc gga agc gga cgc acc aaa atc gcc gtg 96
Val Val Lys Asn Thr Val Ile Gly Ser Gly Arg Thr Lys Ile Ala Val
      20           25           30

ccg ctt gtc gcc cgc gat gcc gcc gta ctt tcc gcc gta ctt gac caa 144
Pro Leu Val Ala Arg Asp Ala Ala Val Leu Ser Ala Val Leu Asp Gln
      35           40           45

atc aaa aac ctg ccc ttc gat atc gtc gag ttc cgc gcc gat ttt ttg 192
Ile Lys Asn Leu Pro Phe Asp Ile Val Glu Phe Arg Ala Asp Phe Leu
      50           55           60

gaa tgc gcg ggc agc atc ggc gaa gta ttg cgc cac acg cag acc gtc 240
Glu Cys Ala Gly Ser Ile Gly Glu Val Leu Arg His Thr Gln Thr Val
      65           70           75           80

cgc gac gcg cta ccc gac aag ccg ctg ctg ttt acg ttc aga cgg cat 288
Arg Asp Ala Leu Pro Asp Lys Pro Leu Leu Phe Thr Phe Arg Arg His
      85           90           95

ggc gaa ggc ggc tcg ttc ccg tgt tcg gac gat tat tat ttt gaa ctg 336
Gly Glu Gly Gly Ser Phe Pro Cys Ser Asp Asp Tyr Tyr Phe Glu Leu
      100          105          110

ctc gac gcg ctg atc gaa agc cgc ctg ccc gac atc atc gac atc gag 384
Leu Asp Ala Leu Ile Glu Ser Arg Leu Pro Asp Ile Ile Asp Ile Glu
      115          120          125

ctg ttt tcc ggc gaa acc gcc gtc cgg tgc gcc gtg gca aat gct caa 432
Leu Phe Ser Gly Glu Thr Ala Val Arg Cys Ala Val Ala Asn Ala Gln
      130          135          140

aaa aac ggc atc gcc gcc ctg ctc tgc aat cat gag ttt cac cgc acg 480
Lys Asn Gly Ile Ala Ala Leu Leu Cys Asn His Glu Phe His Arg Thr

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145              150              155              160

ccg ccg caa gaa gaa atc gta tgc cgt ctg aaa cag atg gag gac tgc 528
Pro Pro Gln Glu Glu Ile Val Cys Arg Leu Lys Gln Met Glu Asp Cys
              165              170              175

ggc gcg gac atc tgc aaa att gcg gtg atg ccg caa agc gcg gaa gat 576
Gly Ala Asp Ile Cys Lys Ile Ala Val Met Pro Gln Ser Ala Glu Asp
              180              185              190

gtg ctg act ttg ctt tcc gcc acg ctc aaa gcg aaa gag ctt gcc gcc 624
Val Leu Thr Leu Leu Ser Ala Thr Leu Lys Ala Lys Glu Leu Ala Ala
              195              200              205

aaa ccg att gtt acg atg tgc atg ggg cag acg ggg gcg gtc agc cgg 672
Lys Pro Ile Val Thr Met Ser Met Gly Gln Thr Gly Ala Val Ser Arg
              210              215              220

ctt gcc gga cag gtg ttc ggc tca agc atc acg ttc ggt tgc gga acg 720
Leu Ala Gly Gln Val Phe Gly Ser Ser Ile Thr Phe Gly Ser Gly Thr
225              230              235              240

caa aac tcc gcg ccg ggg caa atc ggc gta tcc gcc ctc cgt gcg aca 768
Gln Asn Ser Ala Pro Gly Gln Ile Gly Val Ser Ala Leu Arg Ala Thr
              245              250              255

ctc gac tgc ctc gaa aac gga gca gac 795
Leu Asp Cys Leu Glu Asn Gly Ala Asp
              260              265

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<210> 82

<211> 265

<212> PRT

<213> Neisseria meningitidis

<400> 82

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Met Ala Ala Val Phe Ser Val Cys Ala Cys Phe Met Cys Ser Cys Leu
  1              5              10              15

Val Val Lys Asn Thr Val Ile Gly Ser Gly Arg Thr Lys Ile Ala Val
      20              25              30

Pro Leu Val Ala Arg Asp Ala Ala Val Leu Ser Ala Val Leu Asp Gln
      35              40              45

Ile Lys Asn Leu Pro Phe Asp Ile Val Glu Phe Arg Ala Asp Phe Leu
      50              55              60

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Glu Cys Ala Gly Ser Ile Gly Glu Val Leu Arg His Thr Gln Thr Val
 65 70 75 80
 Arg Asp Ala Leu Pro Asp Lys Pro Leu Leu Phe Thr Phe Arg Arg His
 85 90 95
 Gly Glu Gly Gly Ser Phe Pro Cys Ser Asp Asp Tyr Tyr Phe Glu Leu
 100 105 110
 Leu Asp Ala Leu Ile Glu Ser Arg Leu Pro Asp Ile Ile Asp Ile Glu
 115 120 125
 Leu Phe Ser Gly Glu Thr Ala Val Arg Cys Ala Val Ala Asn Ala Gln
 130 135 140
 Lys Asn Gly Ile Ala Ala Leu Leu Cys Asn His Glu Phe His Arg Thr
 145 150 155 160
 Pro Pro Gln Glu Glu Ile Val Cys Arg Leu Lys Gln Met Glu Asp Cys
 165 170 175
 Gly Ala Asp Ile Cys Lys Ile Ala Val Met Pro Gln Ser Ala Glu Asp
 180 185 190
 Val Leu Thr Leu Leu Ser Ala Thr Leu Lys Ala Lys Glu Leu Ala Ala
 195 200 205
 Lys Pro Ile Val Thr Met Ser Met Gly Gln Thr Gly Ala Val Ser Arg
 210 215 220
 Leu Ala Gly Gln Val Phe Gly Ser Ser Ile Thr Phe Gly Ser Gly Thr
 225 230 235 240
 Gln Asn Ser Ala Pro Gly Gln Ile Gly Val Ser Ala Leu Arg Ala Thr
 245 250 255
 Leu Asp Cys Leu Glu Asn Gly Ala Asp
 260 265

<210> 83

<211> 876

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(876)

<400> 83

ttg acc gtc cga acg aaa aag acg gcg cat tat acc cta ttc cat tcc 48
 Leu Thr Val Arg Thr Lys Lys Thr Ala His Tyr Thr Leu Phe His Ser
 1 5 10 15

gac cga aaa ccg aac atg act act ctc aaa ccc gcc ctg ccc gct tat 96
 Asp Arg Lys Pro Asn Met Thr Thr Leu Lys Pro Ala Leu Pro Ala Tyr
 20 25 30

ctg gac aac atc cgc atc atc ctc acg cgc acc agc cat ccc gcc aac 144
 Leu Asp Asn Ile Arg Ile Ile Leu Thr Arg Thr Ser His Pro Ala Asn
 35 40 45

atc ggc tct gcc gcg cgc gcg atg aaa aca atg ggt ctg cac aaa ctg 192
 Ile Gly Ser Ala Ala Arg Ala Met Lys Thr Met Gly Leu His Lys Leu
 50 55 60

acc atc gtc gcc cca aat ctg atg gca acg ccg atg acg gaa aac ccg 240
 Thr Ile Val Ala Pro Asn Leu Met Ala Thr Pro Met Thr Glu Asn Pro
 65 70 75 80

ccc gtg ttt gac ccg gag cat cct caa tcg ttt aaa tta ccg gaa gaa 288
 Pro Val Phe Asp Pro Glu His Pro Gln Ser Phe Lys Leu Pro Glu Glu
 85 90 95

agc ttc atc ctc gct tcc ggc gcg gca gac gtt ttg gaa aat gcc acc 336
 Ser Phe Ile Leu Ala Ser Gly Ala Ala Asp Val Leu Glu Asn Ala Thr
 100 105 110

att gcc gct tct ttg gac gaa gcc ctt gcc gac acc acc atc gcc tgc 384
 Ile Ala Ala Ser Leu Asp Glu Ala Leu Ala Asp Thr Thr Ile Ala Cys
 115 120 125

gcc ctg acc agc cgc cgc cgc gaa att act gcg ccg ctg caa acc ccg 432
 Ala Leu Thr Ser Arg Arg Arg Glu Ile Thr Ala Pro Leu Gln Thr Pro
 130 135 140

cgc gat ttg gta tcc gaa tta ctg cag acc gca aac cga ggc gag aaa 480
 Arg Asp Leu Val Ser Glu Leu Leu Gln Thr Ala Asn Arg Gly Glu Lys
 145 150 155 160

gtg gca ctg gtt ttc ggc aac gag act ttc ggc ttg agc atc gaa gaa 528
 Val Ala Leu Val Phe Gly Asn Glu Thr Phe Gly Leu Ser Ile Glu Glu
 165 170 175

gtc caa gcc tgc aac cga ctg atg acc atc aac ggc aat ccc gac tat 576
 Val Gln Ala Cys Asn Arg Leu Met Thr Ile Asn Gly Asn Pro Asp Tyr
 180 185 190

ttc tcg ctc aac ctc gcc caa gcc gtg cag gtc gtg tgc tac gaa atc 624
 Phe Ser Leu Asn Leu Ala Gln Ala Val Gln Val Val Cys Tyr Glu Ile
 195 200 205

ttc agc caa acc ggt tcg ccc atg acc cat ctt caa caa gaa gac cac 672
 Phe Ser Gln Thr Gly Ser Pro Met Thr His Leu Gln Gln Glu Asp His
 210 215 220

gct gcg acc cac gag caa atc aaa ggc atg gtc gcc cac atg gaa agc 720
 Ala Ala Thr His Glu Gln Ile Lys Gly Met Val Ala His Met Glu Ser
 225 230 235 240

gtg atg aac gac atc ggc ttt ttc aac cgc cgc aac ggc gag cgt ctg 768
 Val Met Asn Asp Ile Gly Phe Phe Asn Arg Arg Asn Gly Glu Arg Leu
 245 250 255

atg cgc cgt atg cag agc ctg ttc gga cgc gcc aac acg caa acc gaa 816
 Met Arg Arg Met Gln Ser Leu Phe Gly Arg Ala Asn Thr Gln Thr Glu
 260 265 270

gac atc gat atc ctg cgc ggt ttt ttc aat acc gtc agc cat cgt atc 864
 Asp Ile Asp Ile Leu Arg Gly Phe Phe Asn Thr Val Ser His Arg Ile
 275 280 285

cat aaa aaa gac 876
 His Lys Lys Asp
 290

<210> 84

<211> 292

<212> PRT

<213> Neisseria meningitidis

<400> 84

Leu Thr Val Arg Thr Lys Lys Thr Ala His Tyr Thr Leu Phe His Ser
 1 5 10 15

Asp Arg Lys Pro Asn Met Thr Thr Leu Lys Pro Ala Leu Pro Ala Tyr
 20 25 30

Leu Asp Asn Ile Arg Ile Ile Leu Thr Arg Thr Ser His Pro Ala Asn
 35 40 45

Ile Gly Ser Ala Ala Arg Ala Met Lys Thr Met Gly Leu His Lys Leu
 50 55 60

Thr Ile Val Ala Pro Asn Leu Met Ala Thr Pro Met Thr Glu Asn Pro
 65 70 75 80

Pro Val Phe Asp Pro Glu His Pro Gln Ser Phe Lys Leu Pro Glu Glu
 85 90 95

Ser Phe Ile Leu Ala Ser Gly Ala Ala Asp Val Leu Glu Asn Ala Thr
 100 105 110

Ile Ala Ala Ser Leu Asp Glu Ala Leu Ala Asp Thr Thr Ile Ala Cys
 115 120 125

Ala Leu Thr Ser Arg Arg Arg Glu Ile Thr Ala Pro Leu Gln Thr Pro
 130 135 140

Arg Asp Leu Val Ser Glu Leu Leu Gln Thr Ala Asn Arg Gly Glu Lys
 145 150 155 160

Val Ala Leu Val Phe Gly Asn Glu Thr Phe Gly Leu Ser Ile Glu Glu
 165 170 175

Val Gln Ala Cys Asn Arg Leu Met Thr Ile Asn Gly Asn Pro Asp Tyr
 180 185 190

Phe Ser Leu Asn Leu Ala Gln Ala Val Gln Val Val Cys Tyr Glu Ile
 195 200 205

Phe Ser Gln Thr Gly Ser Pro Met Thr His Leu Gln Gln Glu Asp His
 210 215 220

Ala Ala Thr His Glu Gln Ile Lys Gly Met Val Ala His Met Glu Ser
 225 230 235 240

Val Met Asn Asp Ile Gly Phe Phe Asn Arg Arg Asn Gly Glu Arg Leu
 245 250 255

Met Arg Arg Met Gln Ser Leu Phe Gly Arg Ala Asn Thr Gln Thr Glu
 260 265 270

Asp Ile Asp Ile Leu Arg Gly Phe Phe Asn Thr Val Ser His Arg Ile
 275 280 285

His Lys Lys Asp
 290

<210> 85

<211> 1545

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(1545)

<400> 85

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gtg cgt ctc aat cat ttc ata atg ata gcg att att ata tat gtg att   48
Val Arg Leu Asn His Phe Ile Met Ile Ala Ile Ile Ile Tyr Val Ile
   1             5             10             15

tcc cct gca aac aag ccg gcc cgc cgc cac agc gtt ccc act tat ccg   96
Ser Pro Ala Asn Lys Pro Ala Arg Arg His Ser Val Pro Thr Tyr Pro
          20             25             30

gct ttg cct tat aat tgc ttt ttt tat gta aca gat tta cct atg aat   144
Ala Leu Pro Tyr Asn Cys Phe Phe Tyr Val Thr Asp Leu Pro Met Asn
          35             40             45

ttc ccc aaa aca gcg gcc tcc ctg ctg ctg ctt ctc gcc tcc ctc gcc   192
Phe Pro Lys Thr Ala Ala Ser Leu Leu Leu Leu Leu Ala Ser Leu Ala
          50             55             60

gca cac gcg ctc gat aca ggt cgc att ccg caa aac gaa atc gcc gta   240
Ala His Ala Leu Asp Thr Gly Arg Ile Pro Gln Asn Glu Ile Ala Val
          65             70             75             80

tat gtc caa gag ctt gac agc gga aaa gtc atc att gac cac cgc tcg   288
Tyr Val Gln Glu Leu Asp Ser Gly Lys Val Ile Ile Asp His Arg Ser
          85             90             95

gat gtc ccc gtc aac ccc gcc tcc aca atg aaa ctc gtt acc gcg ttt   336
Asp Val Pro Val Asn Pro Ala Ser Thr Met Lys Leu Val Thr Ala Phe
          100             105             110

gcc gcc ttc aaa acc ttc ggc agc aat tac cgc tgg gcg acc gag ttt   384
Ala Ala Phe Lys Thr Phe Gly Ser Asn Tyr Arg Trp Ala Thr Glu Phe
          115             120             125

aaa agc aac ggt acg gta aac gac ggc acg ctt gac gga aac ctg tat   432
Lys Ser Asn Gly Thr Val Asn Asp Gly Thr Leu Asp Gly Asn Leu Tyr
          130             135             140

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167

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att ttg acg gac atg aac aag cgt tcg gac aat cta att gcg cgt tcc 1056
Ile Leu Thr Asp Met Asn Lys Arg Ser Asp Asn Leu Ile Ala Arg Ser
      340              345              350

gtc ttc ctc aaa ctc ggc ggc gac ggc aaa ctg ccc gcc gtt tcc gaa 1104
Val Phe Leu Lys Leu Gly Gly Asp Gly Lys Leu Pro Ala Val Ser Glu
      355              360              365

cag gca gcg tct gcc gtc cgg cgt gaa ctt gcc gtg tcg ggc atc gat 1152
Gln Ala Ala Ser Ala Val Arg Arg Glu Leu Ala Val Ser Gly Ile Asp
      370              375              380

gtt gcg gat ttg gtt ttg gaa aac ggt tca ggt ctg tcc aga aaa gaa 1200
Val Ala Asp Leu Val Leu Glu Asn Gly Ser Gly Leu Ser Arg Lys Glu
385              390              395              400

agg gta acg gcg aga atg atg gcg caa atg ttg gaa acg gct tat ttc 1248
Arg Val Thr Ala Arg Met Met Ala Gln Met Leu Glu Thr Ala Tyr Phe
      405              410              415

agc ccg ttt gca caa gat ttc atc gat acg ctg ccc atc gcc ggc aca 1296
Ser Pro Phe Ala Gln Asp Phe Ile Asp Thr Leu Pro Ile Ala Gly Thr
      420              425              430

gac ggg act tta cgc aac cgc ttc aaa caa agc ggc ggg ctg ttg cgc 1344
Asp Gly Thr Leu Arg Asn Arg Phe Lys Gln Ser Gly Gly Leu Leu Arg
      435              440              445

tta aaa acc ggc acg ctc aac aat gtc cgc gcc ctt gca ggt tat tgg 1392
Leu Lys Thr Gly Thr Leu Asn Asn Val Arg Ala Leu Ala Gly Tyr Trp
      450              455              460

ctg ggc gac aaa ccg atg gcg gtg gtc gtc atc atc aac agc ggc cgc 1440
Leu Gly Asp Lys Pro Met Ala Val Val Val Ile Ile Asn Ser Gly Arg
465              470              475              480

gcc gtt tcc ctg ctg ccc gac ttg gac aac ttc gtt gcc aac aac atc 1488
Ala Val Ser Leu Leu Pro Asp Leu Asp Asn Phe Val Ala Asn Asn Ile
      485              490              495

atc tcc ggc ggc gat ggc tgg ctg gat gcg aaa ctg atg tgc aaa gaa 1536
Ile Ser Gly Gly Asp Gly Trp Leu Asp Ala Lys Leu Met Cys Lys Glu
      500              505              510

cgc cga gcc 1545
Arg Arg Ala
      515

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<210> 86

<211> 515

<212> PRT

<213> Neisseria meningitidis

<400> 86

Val Arg Leu Asn His Phe Ile Met Ile Ala Ile Ile Ile Tyr Val Ile
 1 5 10 15

Ser Pro Ala Asn Lys Pro Ala Arg Arg His Ser Val Pro Thr Tyr Pro
 20 25 30

Ala Leu Pro Tyr Asn Cys Phe Phe Tyr Val Thr Asp Leu Pro Met Asn
 35 40 45

Phe Pro Lys Thr Ala Ala Ser Leu Leu Leu Leu Leu Ala Ser Leu Ala
 50 55 60

Ala His Ala Leu Asp Thr Gly Arg Ile Pro Gln Asn Glu Ile Ala Val
 65 70 75 80

Tyr Val Gln Glu Leu Asp Ser Gly Lys Val Ile Ile Asp His Arg Ser
 85 90 95

Asp Val Pro Val Asn Pro Ala Ser Thr Met Lys Leu Val Thr Ala Phe
 100 105 110

Ala Ala Phe Lys Thr Phe Gly Ser Asn Tyr Arg Trp Ala Thr Glu Phe
 115 120 125

Lys Ser Asn Gly Thr Val Asn Asp Gly Thr Leu Asp Gly Asn Leu Tyr
 130 135 140

Trp Ala Gly Ser Gly Asp Pro Val Phe Asn Gln Glu Asn Leu Leu Ala
 145 150 155 160

Val Gln Arg Gln Leu Arg Glu Gln Gly Ile Arg Asn Ile Thr Gly His
 165 170 175

Leu Met Leu Asp His Ser Leu Trp Gly Glu Val Gly Ser Pro Asp Asp
 180 185 190

Phe Glu Ala Asp Ser Gly Ser Pro Phe Met Thr Pro Pro Asn Pro Thr
 195 200 205

Met Leu Ser Ala Gly Met Val Met Val Arg Ala Glu Arg Asn Ala Ala
 210 215 220

Asp Ser Thr Asp Ile Leu Thr Asp Pro Pro Leu Pro His Ile Phe Ala
 225 230 235 240
 Gln Asn Asn Leu Lys Ile Thr Ala Ser Gln Ala Ala Cys Pro Ser Ile
 245 250 255
 Lys Lys Leu Met Arg Ala Ser Phe Ser Asp Asn Thr Leu Lys Leu Arg
 260 265 270
 Gly Asn Ile Pro Glu Ser Cys Leu Gly Lys Pro Val Gly Val Arg Met
 275 280 285
 Phe Ala Leu Asp Glu Leu Ile Arg Gln Ser Phe Thr Asn His Trp Leu
 290 295 300
 Leu Gly Gly Gly Arg Ile Ser Asp Gly Ile Gly Ile Ser Asp Thr Pro
 305 310 315 320
 Glu Gly Ala Gln Thr Leu Ala Val Ala His Ser Lys Pro Met Lys Glu
 325 330 335
 Ile Leu Thr Asp Met Asn Lys Arg Ser Asp Asn Leu Ile Ala Arg Ser
 340 345 350
 Val Phe Leu Lys Leu Gly Gly Asp Gly Lys Leu Pro Ala Val Ser Glu
 355 360 365
 Gln Ala Ala Ser Ala Val Arg Arg Glu Leu Ala Val Ser Gly Ile Asp
 370 375 380
 Val Ala Asp Leu Val Leu Glu Asn Gly Ser Gly Leu Ser Arg Lys Glu
 385 390 395 400
 Arg Val Thr Ala Arg Met Met Ala Gln Met Leu Glu Thr Ala Tyr Phe
 405 410 415
 Ser Pro Phe Ala Gln Asp Phe Ile Asp Thr Leu Pro Ile Ala Gly Thr
 420 425 430
 Asp Gly Thr Leu Arg Asn Arg Phe Lys Gln Ser Gly Gly Leu Leu Arg
 435 440 445
 Leu Lys Thr Gly Thr Leu Asn Asn Val Arg Ala Leu Ala Gly Tyr Trp
 450 455 460
 Leu Gly Asp Lys Pro Met Ala Val Val Val Ile Ile Asn Ser Gly Arg
 465 470 475 480

Ala Val Ser Leu Leu Pro Asp Leu Asp Asn Phe Val Ala Asn Asn Ile
 485 490 495

Ile Ser Gly Gly Asp Gly Trp Leu Asp Ala Lys Leu Met Cys Lys Glu
 500 505 510

Arg Arg Ala
 515

<210> 87

<211> 840

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(840)

<400> 87

atg gat aaa gaa cga att tta acc ccc gca gtc gtg ttt tcc gtc gca 48
 Met Asp Lys Glu Arg Ile Leu Thr Pro Ala Val Val Phe Ser Val Ala
 1 5 10 15

ctg ctg cat ttg gca atg gtg gca ttg ctc tgg cag gcg cac aag ctg 96
 Leu Leu His Leu Ala Met Val Ala Leu Leu Trp Gln Ala His Lys Leu
 20 25 30

ccc gtg ata gag tca ggc aat gtt att gaa ttt gtc gat ttg ggc gat 144
 Pro Val Ile Glu Ser Gly Asn Val Ile Glu Phe Val Asp Leu Gly Asp
 35 40 45

ttt ggc gga ggg gac ggc gca ccc gaa ggt gca ggc gcg cct gcc gcg 192
 Phe Gly Gly Gly Asp Gly Ala Pro Glu Gly Ala Gly Ala Pro Ala Ala
 50 55 60

ccc gaa ccg caa ccc gtg ccc gag ccg ccc aaa cct gtc gag ccg ccc 240
 Pro Glu Pro Gln Pro Val Pro Glu Pro Pro Lys Pro Val Glu Pro Pro
 65 70 75 80

aag ccg gtt ttg aag ccg gtg gtt acg aaa aag gcg gat gcg gat att 288
 Lys Pro Val Leu Lys Pro Val Val Thr Lys Lys Ala Asp Ala Asp Ile
 85 90 95

cag cag cct aag gaa gag ccg aaa cct gaa gaa aag ccg aaa ccc gaa 336
 Gln Gln Pro Lys Glu Glu Pro Lys Pro Glu Glu Lys Pro Lys Pro Glu

100	105	110	
gaa aaa ccg aaa cca gag cct aag ccg gaa gcg aag cct gtc ccg aaa			384
Glu Lys Pro Lys Pro Glu Pro Lys Pro Glu Ala Lys Pro Val Pro Lys			
115	120	125	
ccg gcg gaa aaa ccg gtc gag aag ccg tct gaa aaa cct gcc gaa cat			432
Pro Ala Glu Lys Pro Val Glu Lys Pro Ser Glu Lys Pro Ala Glu His			
130	135	140	
ccc gcc aat gct tct gcc aaa gca gac agc gag cag gcc aat ggg gaa			480
Pro Gly Asn Ala Ser Ala Lys Ala Asp Ser Glu Gln Gly Asn Gly Glu			
145	150	155	160
gat aag gga acc ggt atc aaa gga gac gga acg ggg cgc gga gaa ggc			528
Asp Lys Gly Thr Gly Ile Lys Gly Asp Gly Thr Gly Arg Gly Glu Gly			
165	170	175	
agc ggt aaa ggt agc gcc ggt gtc aaa gcc gaa cac ggg gaa gga gcc			576
Ser Gly Lys Gly Ser Gly Gly Val Lys Gly Glu His Gly Glu Gly Ala			
180	185	190	
ggc agc agc aaa gcc aat cct tta cgc gcc aac gcc agc att ccg cgc			624
Gly Ser Ser Lys Gly Asn Pro Leu Arg Ala Asn Gly Ser Ile Pro Arg			
195	200	205	
ccg gct tat ccc acg ctt tct atg gag aat gat gag cag ggt acg gtt			672
Pro Ala Tyr Pro Thr Leu Ser Met Glu Asn Asp Glu Gln Gly Thr Val			
210	215	220	
gtt ttg agc gtt ttg gtt tct ccg gcc ggt cat gtt gag tcc gtt aaa			720
Val Leu Ser Val Leu Val Ser Pro Gly Gly His Val Glu Ser Val Lys			
225	230	235	240
atc gtg aaa agc agt ggt ttt tcc ccg ttg gac aat gcg gca cgc aag			768
Ile Val Lys Ser Ser Gly Phe Ser Arg Leu Asp Asn Ala Ala Arg Lys			
245	250	255	
gcg gcg caa aac ggg cat ttt caa gcc aat gcc tgg acg gag ttt aaa			816
Ala Ala Gln Asn Gly His Phe Gln Ala Asn Ala Trp Thr Glu Phe Lys			
260	265	270	
gtc ccc gtc aag ttt gaa ttg aat			840
Val Pro Val Lys Phe Glu Leu Asn			
275	280		

<211> 280

<212> PRT

<213> *Neisseria meningitidis*

<400> 88

Met Asp Lys Glu Arg Ile Leu Thr Pro Ala Val Val Phe Ser Val Ala
 1 5 10 15

Leu Leu His Leu Ala Met Val Ala Leu Leu Trp Gln Ala His Lys Leu
 20 25 30

Pro Val Ile Glu Ser Gly Asn Val Ile Glu Phe Val Asp Leu Gly Asp
 35 40 45

Phe Gly Gly Gly Asp Gly Ala Pro Glu Gly Ala Gly Ala Pro Ala Ala
 50 55 60

Pro Glu Pro Gln Pro Val Pro Glu Pro Pro Lys Pro Val Glu Pro Pro
 65 70 75 80

Lys Pro Val Leu Lys Pro Val Val Thr Lys Lys Ala Asp Ala Asp Ile
 85 90 95

Gln Gln Pro Lys Glu Glu Pro Lys Pro Glu Glu Lys Pro Lys Pro Glu
 100 105 110

Glu Lys Pro Lys Pro Glu Pro Lys Pro Glu Ala Lys Pro Val Pro Lys
 115 120 125

Pro Ala Glu Lys Pro Val Glu Lys Pro Ser Glu Lys Pro Ala Glu His
 130 135 140

Pro Gly Asn Ala Ser Ala Lys Ala Asp Ser Glu Gln Gly Asn Gly Glu
 145 150 155 160

Asp Lys Gly Thr Gly Ile Lys Gly Asp Gly Thr Gly Arg Gly Glu Gly
 165 170 175

Ser Gly Lys Gly Ser Gly Gly Val Lys Gly Glu His Gly Glu Gly Ala
 180 185 190

Gly Ser Ser Lys Gly Asn Pro Leu Arg Ala Asn Gly Ser Ile Pro Arg
 195 200 205

Pro Ala Tyr Pro Thr Leu Ser Met Glu Asn Asp Glu Gln Gly Thr Val
 210 215 220

Val Leu Ser Val Leu Val Ser Pro Gly Gly His Val Glu Ser Val Lys

225 230 235 240
 Ile Val Lys Ser Ser Gly Phe Ser Arg Leu Asp Asn Ala Ala Arg Lys
 245 250 255
 Ala Ala Gln Asn Gly His Phe Gln Ala Asn Ala Trp Thr Glu Phe Lys
 260 265 270
 Val Pro Val Lys Phe Glu Leu Asn
 275 280

<210> 89
 <211> 1584
 <212> DNA
 <213> Neisseria meningitidis

<220>
 <221> CDS
 <222> (1)..(1584)

<400> 89
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 Met Ala Leu Phe Leu Ser Ile Phe Pro Ile Val Leu Leu Ile Trp Leu
 1 5 10 15

 atg gtg aaa aaa aac agt atg ccc tcc tac gtc gcg ctg ccg att acc 96
 Met Val Lys Lys Asn Ser Met Pro Ser Tyr Val Ala Leu Pro Ile Thr
 20 25 30

 gca gtg ctg att tac gcc atc aaa ctt ttc tac ttc gac gat gcg ggc 144
 Ala Val Leu Ile Tyr Ala Ile Lys Leu Phe Tyr Phe Asp Asp Ala Gly
 35 40 45

 atg ctg ctc aac gcc acc gcc gct tcc ggc ctc gtc aaa acg ctc acg 192
 Met Leu Leu Asn Ala Thr Ala Ala Ser Gly Leu Val Lys Thr Leu Thr
 50 55 60

 ccg att acc gtg att ttc ggc gcg att atg ttc aac cgt atg atg gaa 240
 Pro Ile Thr Val Ile Phe Gly Ala Ile Met Phe Asn Arg Met Met Glu
 65 70 75 80

 acc acg ggc tgc atc gat gtc atc cgc aaa tgg ctg gcg acc atc agc 288
 Thr Thr Gly Cys Ile Asp Val Ile Arg Lys Trp Leu Ala Thr Ile Ser
 85 90 95

 ccc aac ccc gta gcg caa ctg atg att atc ggc tgg gct ttt gcc ttt 336

Pro Asn Pro Val Ala Gln Leu Met Ile Ile Gly Trp Ala Phe Ala Phe	
100 105 110	
atg att gaa ggc gca tcc ggc ttc ggc acg cct gcc gcg att gcc gcg	384
Met Ile Glu Gly Ala Ser Gly Phe Gly Thr Pro Ala Ala Ile Ala Ala	
115 120 125	
ccg att ctg atg agc ttg ggc ttc aac ccg ttg aaa gtg gcg att ttc	432
Pro Ile Leu Met Ser Leu Gly Phe Asn Pro Leu Lys Val Ala Ile Phe	
130 135 140	
act ttg gtg atg aac tcc gtc ccc gtc tct ttc ggc gcg gtc ggt acg	480
Thr Leu Val Met Asn Ser Val Pro Val Ser Phe Gly Ala Val Gly Thr	
145 150 155 160	
ccg acc tgg ttc ggt ttc gca ccg ctg aac ctg agt gcc gaa gac atc	528
Pro Thr Trp Phe Gly Phe Ala Pro Leu Asn Leu Ser Ala Glu Asp Ile	
165 170 175	
ctc gcc atc ggc agg cag acc ggc gta atg cac ttc ttc gca ggt ttc	576
Leu Ala Ile Gly Arg Gln Thr Gly Val Met His Phe Phe Ala Gly Phe	
180 185 190	
gtc atc ccc gtc atc ggc ttg ggc ttc atc gta cct tgg tct gaa atc	624
Val Ile Pro Val Ile Gly Leu Gly Phe Ile Val Pro Trp Ser Glu Ile	
195 200 205	
cgc aaa aac ttg ggc ttc gtc gcc att gcc gtc ttc tcc tgc acc att	672
Arg Lys Asn Leu Gly Phe Val Ala Ile Ala Val Phe Ser Cys Thr Ile	
210 215 220	
cct tat gtc gca ttg gcg atg gtc aac gaa gaa ttc ccg tcg ctc gtc	720
Pro Tyr Val Ala Leu Ala Met Val Asn Glu Glu Phe Pro Ser Leu Val	
225 230 235 240	
gcc ggc gcg atc ggc ctg atg gtg tcc gta ttc gcc gcc aac caa ggc	768
Ala Gly Ala Ile Gly Leu Met Val Ser Val Phe Ala Ala Asn Gln Gly	
245 250 255	
tgg ggc ttg agc aaa gac cac gcc aaa gac ccg aat gcc gaa aaa gtg	816
Trp Gly Leu Ser Lys Asp His Ala Lys Asp Pro Asn Ala Glu Lys Val	
260 265 270	
ccg ttc gcc caa gtc gcc aaa gca ctc gcc cct ttg ggt atg ctg atc	864
Pro Phe Ala Gln Val Ala Lys Ala Leu Ala Pro Leu Gly Met Leu Ile	
275 280 285	
ggc atg ctg gtg gtt acg cgc atc aaa cag ctc ggc atc aaa ggc att	912

Gly Met Leu Val Val Thr Arg Ile Lys Gln Leu Gly Ile Lys Gly Ile	
290 295 300	
ttg acc agc aaa gaa gaa tgg ttc agc ttc caa ctg ccg ttt gat ttg	960
Leu Thr Ser Lys Glu Glu Trp Phe Ser Phe Gln Leu Pro Phe Asp Leu	
305 310 315 320	
tcc aaa atc acc gtc agc gac tcc ctg acg att act ttc ggc aat att	1008
Ser Lys Ile Thr Val Ser Asp Ser Leu Thr Ile Thr Phe Gly Asn Ile	
325 330 335	
ttc gga caa gat gtc agc gcg tct tac caa acg ctg tac gtc ccg gct	1056
Phe Gly Gln Asp Val Ser Ala Ser Tyr Gln Thr Leu Tyr Val Pro Ala	
340 345 350	
tgg att ccg ttt gtg ctg acc gtt tgg att tgc atc ctg ctg tat aaa	1104
Trp Ile Pro Phe Val Leu Thr Val Trp Ile Cys Ile Leu Leu Tyr Lys	
355 360 365	
acc aaa ttc aaa gat gcc tgg acg att tat tcc gta act ttc aat caa	1152
Thr Lys Phe Lys Asp Ala Trp Thr Ile Tyr Ser Val Thr Phe Asn Gln	
370 375 380	
acc aaa aaa ccg ctg ctt gcc ctg atg ggc gcg ctg att atg gtt cag	1200
Thr Lys Lys Pro Leu Leu Ala Leu Met Gly Ala Leu Ile Met Val Gln	
385 390 395 400	
ctg atg ctg gtc ggc ggc gac aat tcg atg gtg aaa atc atc ggt aag	1248
Leu Met Leu Val Gly Gly Asp Asn Ser Met Val Lys Ile Ile Gly Lys	
405 410 415	
gaa ttt gcc gca atg gcg ggc gaa cac tgg gtt tat ttc tcg ccg tat	1296
Glu Phe Ala Ala Met Ala Gly Glu His Trp Val Tyr Phe Ser Pro Tyr	
420 425 430	
ctg ggc gcg atc ggt gcg ttc ttc tcc ggt tcc aac acc gtg tcc aac	1344
Leu Gly Ala Ile Gly Ala Phe Phe Ser Gly Ser Asn Thr Val Ser Asn	
435 440 445	
ctg acc ttc ggc ccg att cag cag caa atc gcc ttg gat acc ggc ctg	1392
Leu Thr Phe Gly Pro Ile Gln Gln Gln Ile Ala Leu Asp Thr Gly Leu	
450 455 460	
tcc gtt acc ctg att ctg gcg ttg cag tcc gtc ggc ggc gcg atg ggc	1440
Ser Val Thr Leu Ile Leu Ala Leu Gln Ser Val Gly Gly Ala Met Gly	
465 470 475 480	
aat atg gtg tgc ctc aac aac atc atc gcc gta tgt acc gta ttg gat	1488

Asn Met Val Cys Leu Asn Asn Ile Ile Ala Val Cys Thr Val Leu Asp
 485 490 495

gtg aaa aat tcc gaa ggt gcg att atc aag aaa acc gtt atc ccg atg 1536
 Val Lys Asn Ser Glu Gly Ala Ile Ile Lys Lys Thr Val Ile Pro Met
 500 505 510

gcg att tac ggc gtg att gcc gtc gtc gcg gca atg att ttc ttc ctc 1584
 Ala Ile Tyr Gly Val Ile Ala Val Val Ala Ala Met Ile Phe Phe Leu
 515 520 525

<210> 90
 <211> 528
 <212> PRT
 <213> Neisseria meningitidis

<400> 90
 Met Ala Leu Phe Leu Ser Ile Phe Pro Ile Val Leu Leu Ile Trp Leu
 1 5 10 15

Met Val Lys Lys Asn Ser Met Pro Ser Tyr Val Ala Leu Pro Ile Thr
 20 25 30

Ala Val Leu Ile Tyr Ala Ile Lys Leu Phe Tyr Phe Asp Asp Ala Gly
 35 40 45

Met Leu Leu Asn Ala Thr Ala Ala Ser Gly Leu Val Lys Thr Leu Thr
 50 55 60

Pro Ile Thr Val Ile Phe Gly Ala Ile Met Phe Asn Arg Met Met Glu
 65 70 75 80

Thr Thr Gly Cys Ile Asp Val Ile Arg Lys Trp Leu Ala Thr Ile Ser
 85 90 95

Pro Asn Pro Val Ala Gln Leu Met Ile Ile Gly Trp Ala Phe Ala Phe
 100 105 110

Met Ile Glu Gly Ala Ser Gly Phe Gly Thr Pro Ala Ala Ile Ala Ala
 115 120 125

Pro Ile Leu Met Ser Leu Gly Phe Asn Pro Leu Lys Val Ala Ile Phe
 130 135 140

Thr Leu Val Met Asn Ser Val Pro Val Ser Phe Gly Ala Val Gly Thr
 145 150 155 160

Pro Thr Trp Phe Gly Phe Ala Pro Leu Asn Leu Ser Ala Glu Asp Ile
 165 170 175

Leu Ala Ile Gly Arg Gln Thr Gly Val Met His Phe Phe Ala Gly Phe
 180 185 190

Val Ile Pro Val Ile Gly Leu Gly Phe Ile Val Pro Trp Ser Glu Ile
 195 200 205

Arg Lys Asn Leu Gly Phe Val Ala Ile Ala Val Phe Ser Cys Thr Ile
 210 215 220

Pro Tyr Val Ala Leu Ala Met Val Asn Glu Glu Phe Pro Ser Leu Val
 225 230 235 240

Ala Gly Ala Ile Gly Leu Met Val Ser Val Phe Ala Ala Asn Gln Gly
 245 250 255

Trp Gly Leu Ser Lys Asp His Ala Lys Asp Pro Asn Ala Glu Lys Val
 260 265 270

Pro Phe Ala Gln Val Ala Lys Ala Leu Ala Pro Leu Gly Met Leu Ile
 275 280 285

Gly Met Leu Val Val Thr Arg Ile Lys Gln Leu Gly Ile Lys Gly Ile
 290 295 300

Leu Thr Ser Lys Glu Glu Trp Phe Ser Phe Gln Leu Pro Phe Asp Leu
 305 310 315 320

Ser Lys Ile Thr Val Ser Asp Ser Leu Thr Ile Thr Phe Gly Asn Ile
 325 330 335

Phe Gly Gln Asp Val Ser Ala Ser Tyr Gln Thr Leu Tyr Val Pro Ala
 340 345 350

Trp Ile Pro Phe Val Leu Thr Val Trp Ile Cys Ile Leu Leu Tyr Lys
 355 360 365

Thr Lys Phe Lys Asp Ala Trp Thr Ile Tyr Ser Val Thr Phe Asn Gln
 370 375 380

Thr Lys Lys Pro Leu Leu Ala Leu Met Gly Ala Leu Ile Met Val Gln
 385 390 395 400

Leu Met Leu Val Gly Gly Asp Asn Ser Met Val Lys Ile Ile Gly Lys
 405 410 415

Glu Phe Ala Ala Met Ala Gly Glu His Trp Val Tyr Phe Ser Pro Tyr
 420 425 430

Leu Gly Ala Ile Gly Ala Phe Phe Ser Gly Ser Asn Thr Val Ser Asn
 435 440 445

Leu Thr Phe Gly Pro Ile Gln Gln Gln Ile Ala Leu Asp Thr Gly Leu
 450 455 460

Ser Val Thr Leu Ile Leu Ala Leu Gln Ser Val Gly Gly Ala Met Gly
 465 470 475 480

Asn Met Val Cys Leu Asn Asn Ile Ile Ala Val Cys Thr Val Leu Asp
 485 490 495

Val Lys Asn Ser Glu Gly Ala Ile Ile Lys Lys Thr Val Ile Pro Met
 500 505 510

Ala Ile Tyr Gly Val Ile Ala Val Val Ala Ala Met Ile Phe Phe Leu
 515 520 525

<210> 91

<211> 1686

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1686)

<400> 91

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 1 5 10 15

gct tta cca atc atg att tct acc aac ggc atc acc atg cag ttc ggc 96
 Ala Leu Pro Ile Met Ile Ser Thr Asn Gly Ile Thr Met Gln Phe Gly
 20 25 30

gcg aag ccg ctg ttt gaa aac gta tcc gtt aaa ttc ggc gaa ggc aac 144
 Ala Lys Pro Leu Phe Glu Asn Val Ser Val Lys Phe Gly Glu Gly Asn
 35 40 45

cgc tac ggt ttg atc ggc gcg aac ggc tca ggc aaa tcc acc ttc atg 192
 Arg Tyr Gly Leu Ile Gly Ala Asn Gly Ser Gly Lys Ser Thr Phe Met
 50 55 60

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aaa atc ctc ggc ggc gat ttg gaa cag aca gcc ggc gaa gtg gcg att 240
Lys Ile Leu Gly Gly Asp Leu Glu Gln Thr Ala Gly Glu Val Ala Ile
65 70 75 80

gaa aac ggc gtg cgt ttg ggt aaa ttg cgc caa gac cag ttt gcc tac 288
Glu Asn Gly Val Arg Leu Gly Lys Leu Arg Gln Asp Gln Phe Ala Tyr
85 90 95

gaa gat atg cgc gtg ctg gac gtg gta atg atg ggt cac acc gaa atg 336
Glu Asp Met Arg Val Leu Asp Val Val Met Met Gly His Thr Glu Met
100 105 110

tgg gcg gcg atg acc gaa cgt gat gcg att tac gcc aat ccc gaa gcc 384
Trp Ala Ala Met Thr Glu Arg Asp Ala Ile Tyr Ala Asn Pro Glu Ala
115 120 125

acc gaa gac gac tac atg aaa gcc gca gaa ctg gaa gcc aag ttc gcc 432
Thr Glu Asp Asp Tyr Met Lys Ala Ala Glu Leu Glu Ala Lys Phe Ala
130 135 140

gaa tac gac ggc tac acc gcc gaa gcg cgt gcc gcc gag ttg ttg agc 480
Glu Tyr Asp Gly Tyr Thr Ala Glu Ala Arg Ala Ala Glu Leu Leu Ser
145 150 155 160

ggc gtg ggc att tcc gaa gat ttg cac aat gca acc atg gca gaa gtt 528
Gly Val Gly Ile Ser Glu Asp Leu His Asn Ala Thr Met Ala Glu Val
165 170 175

gcc ccg ggc ttc aaa ctg cgc gta ttg ctg gcg caa gcc ctg ttc tcc 576
Ala Pro Gly Phe Lys Leu Arg Val Leu Leu Ala Gln Ala Leu Phe Ser
180 185 190

aag ccg gat gta ttg ctc ttg gac gaa ccg acc aat aac ttg gac att 624
Lys Pro Asp Val Leu Leu Leu Asp Glu Pro Thr Asn Asn Leu Asp Ile
195 200 205

aat acc atc cgt tgg ttg gaa ggc gtg ttg aac cag tac gac tcc acg 672
Asn Thr Ile Arg Trp Leu Glu Gly Val Leu Asn Gln Tyr Asp Ser Thr
210 215 220

atg att atc atc tcg cac gac cgc cac ttc ttg aac gaa gtc tgc acg 720
Met Ile Ile Ile Ser His Asp Arg His Phe Leu Asn Glu Val Cys Thr
225 230 235 240

cat atg gca gat ttg gat tac aac acc atc acc atc tat ccg ggc aac 768
His Met Ala Asp Leu Asp Tyr Asn Thr Ile Thr Ile Tyr Pro Gly Asn
245 250 255

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tac gac gac tat atg ctc gcc tcc gcc caa tcg cgc gaa cgc gcc ctg	816
Tyr Asp Asp Tyr Met Leu Ala Ser Ala Gln Ser Arg Glu Arg Ala Leu	
260 265 270	
aaa gac aat gcc aag gcg aag gaa aaa ttg cag gaa ttg caa gaa ttt	864
Lys Asp Asn Ala Lys Ala Lys Glu Lys Leu Gln Glu Leu Gln Glu Phe	
275 280 285	
gtg gcg cgg ttc agt gcc aac aaa tcc aaa gcc cgt cag gca acc agc	912
Val Ala Arg Phe Ser Ala Asn Lys Ser Lys Ala Arg Gln Ala Thr Ser	
290 295 300	
cgt ctg aaa cag gca gac aaa atc aaa tcg gag atg gtc gag gtc aaa	960
Arg Leu Lys Gln Ala Asp Lys Ile Lys Ser Glu Met Val Glu Val Lys	
305 310 315 320	
ccc tct acc cgt caa aac ccg tat atc cgt ttt gaa gcc gat gaa aaa	1008
Pro Ser Thr Arg Gln Asn Pro Tyr Ile Arg Phe Glu Ala Asp Glu Lys	
325 330 335	
gcc aag ctg cac cgt cag gct gtg gaa gtt gaa aaa ctg gcg aaa cgc	1056
Ala Lys Leu His Arg Gln Ala Val Glu Val Glu Lys Leu Ala Lys Arg	
340 345 350	
ttt gaa acc cag ttg ttt aaa aac ctg aac ttt atc ctt gaa gcg gga	1104
Phe Glu Thr Gln Leu Phe Lys Asn Leu Asn Phe Ile Leu Glu Ala Gly	
355 360 365	
caa cgc ctc gcc atc atc ggc ccg aac ggc gcg ggc aaa tcc acc ctg	1152
Gln Arg Leu Ala Ile Ile Gly Pro Asn Gly Ala Gly Lys Ser Thr Leu	
370 375 380	
ctg aaa ctc ttg gcc ggc gcg tac aac ccc gaa tat tca gac ggt ctg	1200
Leu Lys Leu Leu Ala Gly Ala Tyr Asn Pro Glu Tyr Ser Asp Gly Leu	
385 390 395 400	
ctg ccg gac gaa ggc agc atc aaa tgg gcg gaa aaa gcc agc gtc ggc	1248
Leu Pro Asp Glu Gly Ser Ile Lys Trp Ala Glu Lys Ala Ser Val Gly	
405 410 415	
tac tat ccg caa gat cac gaa aac gac ttc gat gtc gat atg gat ttg	1296
Tyr Tyr Pro Gln Asp His Glu Asn Asp Phe Asp Val Asp Met Asp Leu	
420 425 430	
agc gaa tgg atg cgc caa tgg ggg cag gac ggc gac gac gaa caa gtc	1344
Ser Glu Trp Met Arg Gln Trp Gly Gln Asp Gly Asp Asp Glu Gln Val	
435 440 445	

atc cgc ggc act ttg ggg cgt ttg ctc ttc ggc agt aac gat gtc gtg 1392
 Ile Arg Gly Thr Leu Gly Arg Leu Leu Phe Gly Ser Asn Asp Val Val
 450 455 460

aaa aaa gtg aag gtt ctc tcc ggt ggt gaa aaa ggc cgt atg ctt tac 1440
 Lys Lys Val Lys Val Leu Ser Gly Gly Glu Lys Gly Arg Met Leu Tyr
 465 470 475 480

ggc aaa ctg ttg ctg ttg aaa ccc aat gtc tta gtc atg gat gaa ccg 1488
 Gly Lys Leu Leu Leu Leu Lys Pro Asn Val Leu Val Met Asp Glu Pro
 485 490 495

acc aac cac atg gac atg gaa agc atc gaa tcc ttg aac atg gcg ctg 1536
 Thr Asn His Met Asp Met Glu Ser Ile Glu Ser Leu Asn Met Ala Leu
 500 505 510

gaa aaa tac aac ggc acg ctg att ttc gtc tcc cac gac cgt cag ttc 1584
 Glu Lys Tyr Asn Gly Thr Leu Ile Phe Val Ser His Asp Arg Gln Phe
 515 520 525

gtt tct tcc ctg gct acc caa atc att gaa ttg gac ggc aaa ggc gga 1632
 Val Ser Ser Leu Ala Thr Gln Ile Ile Glu Leu Asp Gly Lys Gly Gly
 530 535 540

tat gaa cac tac ttg ggc gat tac gaa agc tat ctc gag aaa aaa ggc 1680
 Tyr Glu His Tyr Leu Gly Asp Tyr Glu Ser Tyr Leu Glu Lys Lys Gly
 545 550 555 560

gta gca 1686
 Val Ala

<210> 92

<211> 562

<212> PRT

<213> Neisseria meningitidis

<400> 92

Met Leu Gln Ser Glu Asn Ser Lys Asn Leu Ile Ser Cys Ser Ile Lys
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 20 25 30

Ala Lys Pro Leu Phe Glu Asn Val Ser Val Lys Phe Gly Glu Gly Asn
 35 40 45

Arg Tyr Gly Leu Ile Gly Ala Asn Gly Ser Gly Lys Ser Thr Phe Met
 50 55 60

Lys Ile Leu Gly Gly Asp Leu Glu Gln Thr Ala Gly Glu Val Ala Ile
 65 70 75 80

Glu Asn Gly Val Arg Leu Gly Lys Leu Arg Gln Asp Gln Phe Ala Tyr
 85 90 95

Glu Asp Met Arg Val Leu Asp Val Val Met Met Gly His Thr Glu Met
 100 105 110

Trp Ala Ala Met Thr Glu Arg Asp Ala Ile Tyr Ala Asn Pro Glu Ala
 115 120 125

Thr Glu Asp Asp Tyr Met Lys Ala Ala Glu Leu Glu Ala Lys Phe Ala
 130 135 140

Glu Tyr Asp Gly Tyr Thr Ala Glu Ala Arg Ala Ala Glu Leu Leu Ser
 145 150 155 160

Gly Val Gly Ile Ser Glu Asp Leu His Asn Ala Thr Met Ala Glu Val
 165 170 175

Ala Pro Gly Phe Lys Leu Arg Val Leu Leu Ala Gln Ala Leu Phe Ser
 180 185 190

Lys Pro Asp Val Leu Leu Leu Asp Glu Pro Thr Asn Asn Leu Asp Ile
 195 200 205

Asn Thr Ile Arg Trp Leu Glu Gly Val Leu Asn Gln Tyr Asp Ser Thr
 210 215 220

Met Ile Ile Ile Ser His Asp Arg His Phe Leu Asn Glu Val Cys Thr
 225 230 235 240

His Met Ala Asp Leu Asp Tyr Asn Thr Ile Thr Ile Tyr Pro Gly Asn
 245 250 255

Tyr Asp Asp Tyr Met Leu Ala Ser Ala Gln Ser Arg Glu Arg Ala Leu
 260 265 270

Lys Asp Asn Ala Lys Ala Lys Glu Lys Leu Gln Glu Leu Gln Glu Phe
 275 280 285

Val Ala Arg Phe Ser Ala Asn Lys Ser Lys Ala Arg Gln Ala Thr Ser
 290 295 300

Arg Leu Lys Gln Ala Asp Lys Ile Lys Ser Glu Met Val Glu Val Lys
 305 310 315 320
 Pro Ser Thr Arg Gln Asn Pro Tyr Ile Arg Phe Glu Ala Asp Glu Lys
 325 330 335
 Ala Lys Leu His Arg Gln Ala Val Glu Val Glu Lys Leu Ala Lys Arg
 340 345 350
 Phe Glu Thr Gln Leu Phe Lys Asn Leu Asn Phe Ile Leu Glu Ala Gly
 355 360 365
 Gln Arg Leu Ala Ile Ile Gly Pro Asn Gly Ala Gly Lys Ser Thr Leu
 370 375 380
 Leu Lys Leu Leu Ala Gly Ala Tyr Asn Pro Glu Tyr Ser Asp Gly Leu
 385 390 395 400
 Leu Pro Asp Glu Gly Ser Ile Lys Trp Ala Glu Lys Ala Ser Val Gly
 405 410 415
 Tyr Tyr Pro Gln Asp His Glu Asn Asp Phe Asp Val Asp Met Asp Leu
 420 425 430
 Ser Glu Trp Met Arg Gln Trp Gly Gln Asp Gly Asp Asp Glu Gln Val
 435 440 445
 Ile Arg Gly Thr Leu Gly Arg Leu Leu Phe Gly Ser Asn Asp Val Val
 450 455 460
 Lys Lys Val Lys Val Leu Ser Gly Gly Glu Lys Gly Arg Met Leu Tyr
 465 470 475 480
 Gly Lys Leu Leu Leu Lys Pro Asn Val Leu Val Met Asp Glu Pro
 485 490 495
 Thr Asn His Met Asp Met Glu Ser Ile Glu Ser Leu Asn Met Ala Leu
 500 505 510
 Glu Lys Tyr Asn Gly Thr Leu Ile Phe Val Ser His Asp Arg Gln Phe
 515 520 525
 Val Ser Ser Leu Ala Thr Gln Ile Ile Glu Leu Asp Gly Lys Gly Gly
 530 535 540
 Tyr Glu His Tyr Leu Gly Asp Tyr Glu Ser Tyr Leu Glu Lys Lys Gly
 545 550 555 560

Val Ala

<210> 93

<211> 1126

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(1125)

<400> 93

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tac gaa cgc tac cgc tac cgc cgc ctc atc cac gcc gtc cgg ctc ggc 96
Tyr Glu Arg Tyr Arg Tyr Arg Arg Leu Ile His Ala Val Arg Leu Gly
      20             25             30

ggg gcc gtc ctg ttc gcc acc gcc tcc gcc cgg ctg ctc cac ctc caa 144
Gly Ala Val Leu Phe Ala Thr Ala Ser Ala Arg Leu Leu His Leu Gln
      35             40             45

cac gcc gag tgg ata ggg atg acc gtc ttc gtc gtc ctc ggc atg ctc 192
His Gly Glu Trp Ile Gly Met Thr Val Phe Val Val Leu Gly Met Leu
      50             55             60

cag ttt caa ggg gcg att tac tcc aag gcg gtg gaa cgt atg ctc ggc 240
Gln Phe Gln Gly Ala Ile Tyr Ser Lys Ala Val Glu Arg Met Leu Gly
      65             70             75             80

acg gtc atc ggg ctg ggc gcg ggt ttg ggc gtt tta tgg ctg aac cag 288
Thr Val Ile Gly Leu Gly Ala Gly Leu Gly Val Leu Trp Leu Asn Gln
      85             90             95

cat tat ttc cac gcc aac ctc ctc ttc tac ctc acc gtc ggc acg gca 336
His Tyr Phe His Gly Asn Leu Leu Phe Tyr Leu Thr Val Gly Thr Ala
      100            105            110

agc gca ctg gcc ggc tgg gcg gcg gtc ggc aaa aac ggc tac gtc cct 384
Ser Ala Leu Ala Gly Trp Ala Ala Val Gly Lys Asn Gly Tyr Val Pro
      115            120            125

atg ctg gcg ggg ctg acg atg tgc atg ctc atc ggc gac aac ggc agc 432

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Met Leu Ala Gly Leu Thr Met Cys Met Leu Ile Gly Asp Asn Gly Ser	
130 135 140	
gaa tgg ttc gac agc ggc ctg atg cgc gcg atg aac gtc ctc atc ggc	480
Glu Trp Phe Asp Ser Gly Leu Met Arg Ala Met Asn Val Leu Ile Gly	
145 150 155 160	
gcg gcc atc gcc atc gcc gcc gcc aaa ctg ctg ccg ctg aaa tcc aca	528
Ala Ala Ile Ala Ile Ala Ala Ala Lys Leu Leu Pro Leu Lys Ser Thr	
165 170 175	
ctg atg tgg cgt ttc atg ctt gcc gac aac ctg acc gac tgc agc aaa	576
Leu Met Trp Arg Phe Met Leu Ala Asp Asn Leu Thr Asp Cys Ser Lys	
180 185 190	
atg att gcc gaa atc agc aac ggc agg cgc atg acc cgc gaa cgc ctc	624
Met Ile Ala Glu Ile Ser Asn Gly Arg Arg Met Thr Arg Glu Arg Leu	
195 200 205	
gaa gag aac atg gcg aaa atg cgc caa atc aac gca cgc atg gtc aaa	672
Glu Glu Asn Met Ala Lys Met Arg Gln Ile Asn Ala Arg Met Val Lys	
210 215 220	
agc cgc agc cac ctc gcc gcc aca tcg ggc gaa agc cgc atc agc ccc	720
Ser Arg Ser His Leu Ala Ala Thr Ser Gly Glu Ser Arg Ile Ser Pro	
225 230 235 240	
gcc atg atg gaa gcc atg cag cac gcc cac cgt aaa att gtc aac acc	768
Ala Met Met Glu Ala Met Gln His Ala His Arg Lys Ile Val Asn Thr	
245 250 255	
acc gag ctg ctc ctg acc acc gcc gcc aag ctg caa tct ccc aaa ctc	816
Thr Glu Leu Leu Leu Thr Thr Ala Ala Lys Leu Gln Ser Pro Lys Leu	
260 265 270	
aac ggc agc gaa atc cgg ctg ctt gac cgc cac ttc aca ctg ctc caa	864
Asn Gly Ser Glu Ile Arg Leu Leu Asp Arg His Phe Thr Leu Leu Gln	
275 280 285	
acc gac ctg caa caa acc gtc gcc ctt atc aac ggc aga cac gcc cgc	912
Thr Asp Leu Gln Gln Thr Val Ala Leu Ile Asn Gly Arg His Ala Arg	
290 295 300	
cgc atc cgc atc gac acc gcc atc aac ccc gaa ctg gaa gcc ctc gcc	960
Arg Ile Arg Ile Asp Thr Ala Ile Asn Pro Glu Leu Glu Ala Leu Ala	
305 310 315 320	
gaa cac ctc cac tac caa tgg cag ggc ttc ctc tgg ctc agc acc aat	1008

Glu His Leu His Tyr Gln Trp Gln Gly Phe Leu Trp Leu Ser Thr Asn
 325 330 335

atg cgt cag gaa att tcc gcc ctc gtc atc ctg ctg caa cgc acc cgc 1056
 Met Arg Gln Glu Ile Ser Ala Leu Val Ile Leu Leu Gln Arg Thr Arg
 340 345 350

cgc aaa tgg ctg gat gcc cac gaa cgc caa cac ctg cgc caa agc ctg 1104
 Arg Lys Trp Leu Asp Ala His Glu Arg Gln His Leu Arg Gln Ser Leu
 355 360 365

ctt gaa aca cgg gaa cac agt t 1126
 Leu Glu Thr Arg Glu His Ser
 370 375

<210> 94
 <211> 375
 <212> PRT
 <213> Neisseria meningitidis

<400> 94
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Gly Ala Val Leu Phe Ala Thr Ala Ser Ala Arg Leu Leu His Leu Gln
 35 40 45

His Gly Glu Trp Ile Gly Met Thr Val Phe Val Val Leu Gly Met Leu
 50 55 60

Gln Phe Gln Gly Ala Ile Tyr Ser Lys Ala Val Glu Arg Met Leu Gly
 65 70 75 80

Thr Val Ile Gly Leu Gly Ala Gly Leu Gly Val Leu Trp Leu Asn Gln
 85 90 95

His Tyr Phe His Gly Asn Leu Leu Phe Tyr Leu Thr Val Gly Thr Ala
 100 105 110

Ser Ala Leu Ala Gly Trp Ala Ala Val Gly Lys Asn Gly Tyr Val Pro
 115 120 125

Met Leu Ala Gly Leu Thr Met Cys Met Leu Ile Gly Asp Asn Gly Ser
 130 135 140

Glu Trp Phe Asp Ser Gly Leu Met Arg Ala Met Asn Val Leu Ile Gly
 145 150 155 160

Ala Ala Ile Ala Ile Ala Ala Ala Lys Leu Leu Pro Leu Lys Ser Thr
 165 170 175

Leu Met Trp Arg Phe Met Leu Ala Asp Asn Leu Thr Asp Cys Ser Lys
 180 185 190

Met Ile Ala Glu Ile Ser Asn Gly Arg Arg Met Thr Arg Glu Arg Leu
 195 200 205

Glu Glu Asn Met Ala Lys Met Arg Gln Ile Asn Ala Arg Met Val Lys
 210 215 220

Ser Arg Ser His Leu Ala Ala Thr Ser Gly Glu Ser Arg Ile Ser Pro
 225 230 235 240

Ala Met Met Glu Ala Met Gln His Ala His Arg Lys Ile Val Asn Thr
 245 250 255

Thr Glu Leu Leu Leu Thr Thr Ala Ala Lys Leu Gln Ser Pro Lys Leu
 260 265 270

Asn Gly Ser Glu Ile Arg Leu Leu Asp Arg His Phe Thr Leu Leu Gln
 275 280 285

Thr Asp Leu Gln Gln Thr Val Ala Leu Ile Asn Gly Arg His Ala Arg
 290 295 300

Arg Ile Arg Ile Asp Thr Ala Ile Asn Pro Glu Leu Glu Ala Leu Ala
 305 310 315 320

Glu His Leu His Tyr Gln Trp Gln Gly Phe Leu Trp Leu Ser Thr Asn
 325 330 335

Met Arg Gln Glu Ile Ser Ala Leu Val Ile Leu Leu Gln Arg Thr Arg
 340 345 350

Arg Lys Trp Leu Asp Ala His Glu Arg Gln His Leu Arg Gln Ser Leu
 355 360 365

Leu Glu Thr Arg Glu His Ser
 370 375

<210> 95

<211> 1725

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(1725)

<400> 95

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  1             5             10            15

ggg gtc gag tac gtt ttc ggt tat ccc ggc ggt gcg gtt atc gaa atc 96
Gly Val Glu Tyr Val Phe Gly Tyr Pro Gly Gly Ala Val Ile Glu Ile
      20             25             30

tac gat gcc ctt ttc caa ctc aat aaa ttc aag cac att ctg acc cgt 144
Tyr Asp Ala Leu Phe Gln Leu Asn Lys Phe Lys His Ile Leu Thr Arg
      35             40             45

cac gag cag gcg gca gta cac gcg gca gat gcg tat gcg cgc gtc agc 192
His Glu Gln Ala Ala Val His Ala Ala Asp Ala Tyr Ala Arg Val Ser
      50             55             60

ggg aag gtg ggc gtg gca ttg gtt aca tcc ggc ccg ggc gtt acc aat 240
Gly Lys Val Gly Val Ala Leu Val Thr Ser Gly Pro Gly Val Thr Asn
      65             70             75             80

gca ctg acc ggt att gct act gcc tat acg gat tcg att ccg atg gtg 288
Ala Leu Thr Gly Ile Ala Thr Ala Tyr Thr Asp Ser Ile Pro Met Val
      85             90             95

gtc atc agc ggg cag gta ggc aat tcc ctg att ggt acg gat gcg ttc 336
Val Ile Ser Gly Gln Val Gly Asn Ser Leu Ile Gly Thr Asp Ala Phe
      100            105            110

caa gaa gtt gat acg gtg ggt att acc cgt ccg tgc gtc aaa cac aat 384
Gln Glu Val Asp Thr Val Gly Ile Thr Arg Pro Cys Val Lys His Asn
      115            120            125

ttc ctg gtt acg gac atc aat gag ttg gcg gaa acc att aaa aag gcg 432
Phe Leu Val Thr Asp Ile Asn Glu Leu Ala Glu Thr Ile Lys Lys Ala
      130            135            140

ttc caa att gcc gca agc ggc cga ccg ggg ccc gtg gtg gtt gat gtc 480
Phe Gln Ile Ala Ala Ser Gly Arg Pro Gly Pro Val Val Val Asp Val

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145	150	155	160	
ccg aaa gat gtt acg cag gcg atg gcg aaa ttc agc tat cct cag gaa				528
Pro Lys Asp Val Thr Gln Ala Met Ala Lys Phe Ser Tyr Pro Gln Glu				
	165	170	175	
gac att ttt atc cgt tcg tat caa ccg gtt gtt caa ggg cat atc ggt				576
Asp Ile Phe Ile Arg Ser Tyr Gln Pro Val Val Gln Gly His Ile Gly				
	180	185	190	
cag att aaa aag gcc gtg cag atg ttg gca tct gcc aaa cgt ccg gtc				624
Gln Ile Lys Lys Ala Val Gln Met Leu Ala Ser Ala Lys Arg Pro Val				
	195	200	205	
gtc tat ttt ggc ggc ggc gtg gtg ttg ggt aat gct tct gaa gag ctg				672
Val Tyr Phe Gly Gly Gly Val Val Leu Gly Asn Ala Ser Glu Glu Leu				
	210	215	220	
acc cga ttt gtc cga atg acg ggt gct ccg tgc acg ggt acg ctg atg				720
Thr Arg Phe Val Arg Met Thr Gly Ala Pro Cys Thr Gly Thr Leu Met				
	225	230	235	240
ggc ttg ggc gct tat cct tcc ggc gac cgc caa ttc ctc ggt atg ctc				768
Gly Leu Gly Ala Tyr Pro Ser Gly Asp Arg Gln Phe Leu Gly Met Leu				
	245	250	255	
ggt atg cac ggt act tac gag gca aac ctt gcc atg cag aat gcg gat				816
Gly Met His Gly Thr Tyr Glu Ala Asn Leu Ala Met Gln Asn Ala Asp				
	260	265	270	
gtc gtt ctt gcc gta ggt gcg cgt ttt gac gac cgt gtg gtt tcc gta				864
Val Val Leu Ala Val Gly Ala Arg Phe Asp Asp Arg Val Val Ser Val				
	275	280	285	
ccg tcc aaa ttt ttc gag aag gcg aag aag gtt atc cat atc gat gtc				912
Pro Ser Lys Phe Phe Glu Lys Ala Lys Lys Val Ile His Ile Asp Val				
	290	295	300	
gat cct tcc agc atc gcc aaa cgc gtg aag gcg gac att ccg att gtc				960
Asp Pro Ser Ser Ile Ala Lys Arg Val Lys Ala Asp Ile Pro Ile Val				
	305	310	315	320
ggc gac gtg aaa aac gtt ttg tcc gag atg gtt gcg ctg tgg caa aaa				1008
Gly Asp Val Lys Asn Val Leu Ser Glu Met Val Ala Leu Trp Gln Lys				
	325	330	335	
caa gag tcc gtg ccg tct gaa gat gct ttg ggc aaa tgg tgg aaa acc				1056
Gln Glu Ser Val Pro Ser Glu Asp Ala Leu Gly Lys Trp Trp Lys Thr				

340	345	350	
ata gag gaa tgg cgt tcc cgc gat tgc ttg tgg ttt gac aac ggc agc			1104
Ile Glu Glu Trp Arg Ser Arg Asp Cys Leu Trp Phe Asp Asn Gly Ser			
355	360	365	
gaa att atc aag cca caa tat gtg att cag aag ctt gcc gag att acc			1152
Glu Ile Ile Lys Pro Gln Tyr Val Ile Gln Lys Leu Ala Glu Ile Thr			
370	375	380	
ggc aat tcg gca atc atc aca tcg gat gta ggg cag cat caa atg ttt			1200
Gly Asn Ser Ala Ile Ile Thr Ser Asp Val Gly Gln His Gln Met Phe			
385	390	395	400
gcg gct caa tat tat ccc ttc gaa cgt ccg cgc caa tgg ctc aac tcc			1248
Ala Ala Gln Tyr Tyr Pro Phe Glu Arg Pro Arg Gln Trp Leu Asn Ser			
405	410	415	
ggc ggt ttg ggt acg atg ggc gtt ggt ctg cct tat gcg att ggt gca			1296
Gly Gly Leu Gly Thr Met Gly Val Gly Leu Pro Tyr Ala Ile Gly Ala			
420	425	430	
aaa ctt gcc gcc ccg gat caa gac gta ttc tgt att acc ggc gac ggc			1344
Lys Leu Ala Ala Pro Asp Gln Asp Val Phe Cys Ile Thr Gly Asp Gly			
435	440	445	
tcg atc cag atg aac atc caa gag ctg tcc acc tgt ttc caa tat cgg			1392
Ser Ile Gln Met Asn Ile Gln Glu Leu Ser Thr Cys Phe Gln Tyr Arg			
450	455	460	
att ccg gtt aac gtc att acg ctg aac aac ggt tat ctc ggt atg gta			1440
Ile Pro Val Asn Val Ile Thr Leu Asn Asn Gly Tyr Leu Gly Met Val			
465	470	475	480
cgc cag tgg cag gaa ata tat tac ggc gga cga gag tcg gaa acc tat			1488
Arg Gln Trp Gln Glu Ile Tyr Tyr Gly Gly Arg Glu Ser Glu Thr Tyr			
485	490	495	
ttc gat tct ttg ccc gat ttc gtc aaa ctt gcc gag gca tac ggc cat			1536
Phe Asp Ser Leu Pro Asp Phe Val Lys Leu Ala Glu Ala Tyr Gly His			
500	505	510	
atc ggt atc cgc gtg gac aag aag tct gat gtg gaa ggt gcg ttg ttg			1584
Ile Gly Ile Arg Val Asp Lys Lys Ser Asp Val Glu Gly Ala Leu Leu			
515	520	525	
gaa gca ttg aac caa aaa gac agg ctg gtg ttt atc gac ttc ctg acc			1632
Glu Ala Leu Asn Gln Lys Asp Arg Leu Val Phe Ile Asp Phe Leu Thr			

530 535 540
 gac cag aaa cag aat gtg atg ccc atg gtc ggc aac ggc aaa ggt ttg 1680
 Asp Gln Lys Gln Asn Val Met Pro Met Val Gly Asn Gly Lys Gly Leu
 545 550 555 560
 gac gaa atg gta ctt ccg ccg cat atg cgt gcg gac gga aag gcg 1725
 Asp Glu Met Val Leu Pro Pro His Met Arg Ala Asp Gly Lys Ala
 565 570 575

 <210> 96
 <211> 575
 <212> PRT
 <213> Neisseria meningitidis

 <400> 96
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 Gly Val Glu Tyr Val Phe Gly Tyr Pro Gly Gly Ala Val Ile Glu Ile
 20 25 30
 Tyr Asp Ala Leu Phe Gln Leu Asn Lys Phe Lys His Ile Leu Thr Arg
 35 40 45
 His Glu Gln Ala Ala Val His Ala Ala Asp Ala Tyr Ala Arg Val Ser
 50 55 60
 Gly Lys Val Gly Val Ala Leu Val Thr Ser Gly Pro Gly Val Thr Asn
 65 70 75 80
 Ala Leu Thr Gly Ile Ala Thr Ala Tyr Thr Asp Ser Ile Pro Met Val
 85 90 95
 Val Ile Ser Gly Gln Val Gly Asn Ser Leu Ile Gly Thr Asp Ala Phe
 100 105 110
 Gln Glu Val Asp Thr Val Gly Ile Thr Arg Pro Cys Val Lys His Asn
 115 120 125
 Phe Leu Val Thr Asp Ile Asn Glu Leu Ala Glu Thr Ile Lys Lys Ala
 130 135 140
 Phe Gln Ile Ala Ala Ser Gly Arg Pro Gly Pro Val Val Val Asp Val
 145 150 155 160
 Pro Lys Asp Val Thr Gln Ala Met Ala Lys Phe Ser Tyr Pro Gln Glu

	165		170		175
Asp Ile Phe Ile Arg Ser Tyr Gln Pro Val Val Gln Gly His Ile Gly					
	180		185		190
Gln Ile Lys Lys Ala Val Gln Met Leu Ala Ser Ala Lys Arg Pro Val					
	195		200		205
Val Tyr Phe Gly Gly Gly Val Val Leu Gly Asn Ala Ser Glu Glu Leu					
	210		215		220
Thr Arg Phe Val Arg Met Thr Gly Ala Pro Cys Thr Gly Thr Leu Met					
	225		230		235
					240
Gly Leu Gly Ala Tyr Pro Ser Gly Asp Arg Gln Phe Leu Gly Met Leu					
	245		250		255
Gly Met His Gly Thr Tyr Glu Ala Asn Leu Ala Met Gln Asn Ala Asp					
	260		265		270
Val Val Leu Ala Val Gly Ala Arg Phe Asp Asp Arg Val Val Ser Val					
	275		280		285
Pro Ser Lys Phe Phe Glu Lys Ala Lys Lys Val Ile His Ile Asp Val					
	290		295		300
Asp Pro Ser Ser Ile Ala Lys Arg Val Lys Ala Asp Ile Pro Ile Val					
	305		310		315
					320
Gly Asp Val Lys Asn Val Leu Ser Glu Met Val Ala Leu Trp Gln Lys					
	325		330		335
Gln Glu Ser Val Pro Ser Glu Asp Ala Leu Gly Lys Trp Trp Lys Thr					
	340		345		350
Ile Glu Glu Trp Arg Ser Arg Asp Cys Leu Trp Phe Asp Asn Gly Ser					
	355		360		365
Glu Ile Ile Lys Pro Gln Tyr Val Ile Gln Lys Leu Ala Glu Ile Thr					
	370		375		380
Gly Asn Ser Ala Ile Ile Thr Ser Asp Val Gly Gln His Gln Met Phe					
	385		390		395
					400
Ala Ala Gln Tyr Tyr Pro Phe Glu Arg Pro Arg Gln Trp Leu Asn Ser					
	405		410		415
Gly Gly Leu Gly Thr Met Gly Val Gly Leu Pro Tyr Ala Ile Gly Ala					

420 425 430
 Lys Leu Ala Ala Pro Asp Gln Asp Val Phe Cys Ile Thr Gly Asp Gly
 435 440 445
 Ser Ile Gln Met Asn Ile Gln Glu Leu Ser Thr Cys Phe Gln Tyr Arg
 450 455 460
 Ile Pro Val Asn Val Ile Thr Leu Asn Asn Gly Tyr Leu Gly Met Val
 465 470 475 480
 Arg Gln Trp Gln Glu Ile Tyr Tyr Gly Gly Arg Glu Ser Glu Thr Tyr
 485 490 495
 Phe Asp Ser Leu Pro Asp Phe Val Lys Leu Ala Glu Ala Tyr Gly His
 500 505 510
 Ile Gly Ile Arg Val Asp Lys Lys Ser Asp Val Glu Gly Ala Leu Leu
 515 520 525
 Glu Ala Leu Asn Gln Lys Asp Arg Leu Val Phe Ile Asp Phe Leu Thr
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 Asp Gln Lys Gln Asn Val Met Pro Met Val Gly Asn Gly Lys Gly Leu
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 Asp Glu Met Val Leu Pro Pro His Met Arg Ala Asp Gly Lys Ala
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<210> 97

<211> 570

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(570)

<400> 97

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 1 5 10 15

ata cgc cat acc ccg tcg ccc aat ttc agc ccg agg gaa acg ggg gaa 96
 Ile Arg His Thr Pro Ser Pro Asn Phe Ser Pro Arg Glu Thr Gly Glu
 20 25 30

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acg gtt tcc ctg atc gtg ttg cac aac att tca ctg ccg ccg ttc gaa 144
Thr Val Ser Leu Ile Val Leu His Asn Ile Ser Leu Pro Pro Phe Glu
      35              40              45

tac ggc acg gat gct gtg gaa aag ctg ttt gcc aac cgg ctc gac ccc 192
Tyr Gly Thr Asp Ala Val Glu Lys Leu Phe Ala Asn Arg Leu Asp Pro
      50              55              60

aac gga cat ccg ttc ttc agc ctg ata cac act ttg cgc gta tcc agc 240
Asn Gly His Pro Phe Phe Ser Leu Ile His Thr Leu Arg Val Ser Ser
      65              70              75              80

cat ttc tta atc aaa cgc gac ggc aaa acg gtg cag ttc gta tca tgc 288
His Phe Leu Ile Lys Arg Asp Gly Lys Thr Val Gln Phe Val Ser Cys
      85              90              95

ggc gat atg gcg tac cac gcg ggc gta tcc tcg ttt cgc gga cgg gaa 336
Gly Asp Met Ala Tyr His Ala Gly Val Ser Ser Phe Arg Gly Arg Glu
      100              105              110

aaa tgc aac gca ttt tcc atc ggc atc gaa ttg gaa ggc tgc gat ttc 384
Lys Cys Asn Ala Phe Ser Ile Gly Ile Glu Leu Glu Gly Cys Asp Phe
      115              120              125

gaa ccc ttt acc gaa gcg caa tac cgt tcg ctc gaa aca ttg ttg gaa 432
Glu Pro Phe Thr Glu Ala Gln Tyr Arg Ser Leu Glu Thr Leu Leu Glu
      130              135              140

gca ctc tgc cgc cgc tac ccc gtt acc gca gta acc gga cat cag gac 480
Ala Leu Cys Arg Arg Tyr Pro Val Thr Ala Val Thr Gly His Gln Asp
      145              150              155              160

atc gcg ccc ggc cgc aaa acc gac ccc ggc cac ttt ttc gac tgg cgg 528
Ile Ala Pro Gly Arg Lys Thr Asp Pro Gly His Phe Phe Asp Trp Arg
      165              170              175

cgg ata cgg gag aaa ggg ttt ccc gta gac aga aat gcc gtc 570
Arg Ile Arg Glu Lys Gly Phe Pro Val Asp Arg Asn Ala Val
      180              185              190

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<210> 98

<211> 190

<212> PRT

<213> Neisseria meningitidis

<400> 98

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 35 40 45
 Tyr Gly Thr Asp Ala Val Glu Lys Leu Phe Ala Asn Arg Leu Asp Pro
 50 55 60
 Asn Gly His Pro Phe Phe Ser Leu Ile His Thr Leu Arg Val Ser Ser
 65 70 75 80
 His Phe Leu Ile Lys Arg Asp Gly Lys Thr Val Gln Phe Val Ser Cys
 85 90 95
 Gly Asp Met Ala Tyr His Ala Gly Val Ser Ser Phe Arg Gly Arg Glu
 100 105 110
 Lys Cys Asn Ala Phe Ser Ile Gly Ile Glu Leu Glu Gly Cys Asp Phe
 115 120 125
 Glu Pro Phe Thr Glu Ala Gln Tyr Arg Ser Leu Glu Thr Leu Leu Glu
 130 135 140
 Ala Leu Cys Arg Arg Tyr Pro Val Thr Ala Val Thr Gly His Gln Asp
 145 150 155 160
 Ile Ala Pro Gly Arg Lys Thr Asp Pro Gly His Phe Phe Asp Trp Arg
 165 170 175
 Arg Ile Arg Glu Lys Gly Phe Pro Val Asp Arg Asn Ala Val
 180 185 190

<210> 99

<211> 1515

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1515)

<400> 99

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gtt tct gcc gcc tat ggg gcg gat gcg ccc gcg att ttg gat gac aag	96
Val Ser Ala Ala Tyr Gly Ala Asp Ala Pro Ala Ile Leu Asp Asp Lys	
20 25 30	
gca ttg ttg cag gtg cag cgg tcg gtg tcg gat aag tgg gcg gaa tcg	144
Ala Leu Leu Gln Val Gln Arg Ser Val Ser Asp Lys Trp Ala Glu Ser	
35 40 45	
gat tgg aaa gtt gac aat gat gcc ccg cgc gtg gtt gac ggg gat ttt	192
Asp Trp Lys Val Asp Asn Asp Ala Pro Arg Val Val Asp Gly Asp Phe	
50 55 60	
ttg ttg gcg cat ccg aaa atg ttg gaa cat agt ttg cgc gac gtg ctc	240
Leu Leu Ala His Pro Lys Met Leu Glu His Ser Leu Arg Asp Val Leu	
65 70 75 80	
aac ggc aat cag gcg gat ttg atc gct tcg ttg gcg gat ttg tat gcc	288
Asn Gly Asn Gln Ala Asp Leu Ile Ala Ser Leu Ala Asp Leu Tyr Ala	
85 90 95	
aag ctg ccg gat tat gac gcg gtt ttg tac ggc agg gcg cgg gct ttg	336
Lys Leu Pro Asp Tyr Asp Ala Val Leu Tyr Gly Arg Ala Arg Ala Leu	
100 105 110	
ctg gcg aaa ttg gcg gga agg ccg gcg gag gcg gtg gcg cgg tat cgg	384
Leu Ala Lys Leu Ala Gly Arg Pro Ala Glu Ala Val Ala Arg Tyr Arg	
115 120 125	
gaa ctg cac ggg gaa aat gcg gca gac gag cgg att ttg ctg gat ttg	432
Glu Leu His Gly Glu Asn Ala Ala Asp Glu Arg Ile Leu Leu Asp Leu	
130 135 140	
gcg gcg gcg gag ttt gac gat ttc ccg ctg aag tcg gca gaa agg cat	480
Ala Ala Ala Glu Phe Asp Asp Phe Arg Leu Lys Ser Ala Glu Arg His	
145 150 155 160	
ttt gcc gag gcg gaa aaa ttg gat ttg ccg gcg ccg gtt ttg gaa aat	528
Phe Ala Glu Ala Glu Lys Leu Asp Leu Pro Ala Pro Val Leu Glu Asn	
165 170 175	
gtg ggg cgt ttt ccg aaa aaa gcg gag ggg ctg acg ggc tgg cgt ttt	576
Val Gly Arg Phe Arg Lys Lys Ala Glu Gly Leu Thr Gly Trp Arg Phe	
180 185 190	

tcg ggc ggc atc agt ccg gcg gtc aat aga aat gcc aat aat gcc gcg	624
Ser Gly Gly Ile Ser Pro Ala Val Asn Arg Asn Ala Asn Asn Ala Ala	
195 200 205	
ccg cag tat tgc cgg caa aac gga ggc cgg cag ata tgc agt gtc agc	672
Pro Gln Tyr Cys Arg Gln Asn Gly Gly Arg Gln Ile Cys Ser Val Ser	
210 215 220	
cgg gcg gag cgg gcg gca ggc ttg aat tat gaa atc gag gcg gaa aaa	720
Arg Ala Glu Arg Ala Ala Gly Leu Asn Tyr Glu Ile Glu Ala Glu Lys	
225 230 235 240	
ctg acg gcg ttg gca gat aat cat tat ttg ttg ttc cgt tcc aat atc	768
Leu Thr Ala Leu Ala Asp Asn His Tyr Leu Leu Phe Arg Ser Asn Ile	
245 250 255	
ggc ggc acg agc tat tat ttc agt aaa aaa tca gct tat gac gac ggg	816
Gly Gly Thr Ser Tyr Tyr Phe Ser Lys Lys Ser Ala Tyr Asp Asp Gly	
260 265 270	
ttc ggc aga gcg tat ttg ggt tgg cag tat aaa aat gca cgg cag acg	864
Phe Gly Arg Ala Tyr Leu Gly Trp Gln Tyr Lys Asn Ala Arg Gln Thr	
275 280 285	
gcg ggg att ttg ccg ttt tat cag gtg cag ttg tcg ggc agc gac ggc	912
Ala Gly Ile Leu Pro Phe Tyr Gln Val Gln Leu Ser Gly Ser Asp Gly	
290 295 300	
ttt gat gcg aaa aca aaa cgg gta aac aac cgc cgc ctg ccg ccg tat	960
Phe Asp Ala Lys Thr Lys Arg Val Asn Asn Arg Arg Leu Pro Pro Tyr	
305 310 315 320	
atg ctg gcg cac gga gtc ggc gtg cag ttg tcc cat act tac cgc cca	1008
Met Leu Ala His Gly Val Gly Val Gln Leu Ser His Thr Tyr Arg Pro	
325 330 335	
aac ccg gga tgg caa ttt tcg gtc gcg ctg gaa cat tac cgc caa cgc	1056
Asn Pro Gly Trp Gln Phe Ser Val Ala Leu Glu His Tyr Arg Gln Arg	
340 345 350	
tac cgc gaa cag gat agg gcg gaa tac aat aac ggt cgg cag gac ggg	1104
Tyr Arg Glu Gln Asp Arg Ala Glu Tyr Asn Asn Gly Arg Gln Asp Gly	
355 360 365	
ttt tat gtt tcg tcg gca aaa cgt ttg ggc gaa tcg gca act gtg ttc	1152
Phe Tyr Val Ser Ser Ala Lys Arg Leu Gly Glu Ser Ala Thr Val Phe	
370 375 380	

ggc ggc tgg cag ttt gtg cgg ttt gtg ccg aaa cgc gaa acg gtg ggc 1200
 Gly Gly Trp Gln Phe Val Arg Phe Val Pro Lys Arg Glu Thr Val Gly
 385 390 395 400

ggc gcg gtc aat aat gcc gcc tac cgg cgc aac ggt gtt tat gcc ggc 1248
 Gly Ala Val Asn Asn Ala Ala Tyr Arg Arg Asn Gly Val Tyr Ala Gly
 405 410 415

tgg gcg cag gag tgg cgg cag ttg ggc ggt ttg aac agt cgg gtt tcc 1296
 Trp Ala Gln Glu Trp Arg Gln Leu Gly Gly Leu Asn Ser Arg Val Ser
 420 425 430

gcg tct tat gcc cgc cgc aac tat aag ggc gtt gcg gct ttc tcg aca 1344
 Ala Ser Tyr Ala Arg Arg Asn Tyr Lys Gly Val Ala Ala Phe Ser Thr
 435 440 445

gag gcg caa cgc aac cgc gaa tgg aat gtc tcg ctg gct ttg agc cac 1392
 Glu Ala Gln Arg Asn Arg Glu Trp Asn Val Ser Leu Ala Leu Ser His
 450 455 460

gac aag ttg tcg tac aaa ggt atc gtg ccc gcg ttg aat tat cgt ttc 1440
 Asp Lys Leu Ser Tyr Lys Gly Ile Val Pro Ala Leu Asn Tyr Arg Phe
 465 470 475 480

ggc agg acg gaa agt aat gtg ccg tat gcg aaa cgc cgc aac agc gag 1488
 Gly Arg Thr Glu Ser Asn Val Pro Tyr Ala Lys Arg Arg Asn Ser Glu
 485 490 495

gtg ttt gtg tcg gcg gat tgg cgg ttt 1515
 Val Phe Val Ser Ala Asp Trp Arg Phe
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<210> 100

<211> 505

<212> PRT

<213> Neisseria meningitidis

<400> 100

Met Leu Tyr Phe Arg Tyr Gly Phe Leu Val Val Trp Cys Ala Ala Gly
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Val Ser Ala Ala Tyr Gly Ala Asp Ala Pro Ala Ile Leu Asp Asp Lys
 20 25 30

Ala Leu Leu Gln Val Gln Arg Ser Val Ser Asp Lys Trp Ala Glu Ser
 35 40 45

Asp Trp Lys Val Asp Asn Asp Ala Pro Arg Val Val Asp Gly Asp Phe
 50 55 60

Leu Leu Ala His Pro Lys Met Leu Glu His Ser Leu Arg Asp Val Leu
 65 70 75 80

Asn Gly Asn Gln Ala Asp Leu Ile Ala Ser Leu Ala Asp Leu Tyr Ala
 85 90 95

Lys Leu Pro Asp Tyr Asp Ala Val Leu Tyr Gly Arg Ala Arg Ala Leu
 100 105 110

Leu Ala Lys Leu Ala Gly Arg Pro Ala Glu Ala Val Ala Arg Tyr Arg
 115 120 125

Glu Leu His Gly Glu Asn Ala Ala Asp Glu Arg Ile Leu Leu Asp Leu
 130 135 140

Ala Ala Ala Glu Phe Asp Asp Phe Arg Leu Lys Ser Ala Glu Arg His
 145 150 155 160

Phe Ala Glu Ala Glu Lys Leu Asp Leu Pro Ala Pro Val Leu Glu Asn
 165 170 175

Val Gly Arg Phe Arg Lys Lys Ala Glu Gly Leu Thr Gly Trp Arg Phe
 180 185 190

Ser Gly Gly Ile Ser Pro Ala Val Asn Arg Asn Ala Asn Asn Ala Ala
 195 200 205

Pro Gln Tyr Cys Arg Gln Asn Gly Gly Arg Gln Ile Cys Ser Val Ser
 210 215 220

Arg Ala Glu Arg Ala Ala Gly Leu Asn Tyr Glu Ile Glu Ala Glu Lys
 225 230 235 240

Leu Thr Ala Leu Ala Asp Asn His Tyr Leu Leu Phe Arg Ser Asn Ile
 245 250 255

Gly Gly Thr Ser Tyr Tyr Phe Ser Lys Lys Ser Ala Tyr Asp Asp Gly
 260 265 270

Phe Gly Arg Ala Tyr Leu Gly Trp Gln Tyr Lys Asn Ala Arg Gln Thr
 275 280 285

Ala Gly Ile Leu Pro Phe Tyr Gln Val Gln Leu Ser Gly Ser Asp Gly
 290 295 300

Phe Asp Ala Lys Thr Lys Arg Val Asn Asn Arg Arg Leu Pro Pro Tyr
 305 310 315 320

Met Leu Ala His Gly Val Gly Val Gln Leu Ser His Thr Tyr Arg Pro
 325 330 335

Asn Pro Gly Trp Gln Phe Ser Val Ala Leu Glu His Tyr Arg Gln Arg
 340 345 350

Tyr Arg Glu Gln Asp Arg Ala Glu Tyr Asn Asn Gly Arg Gln Asp Gly
 355 360 365

Phe Tyr Val Ser Ser Ala Lys Arg Leu Gly Glu Ser Ala Thr Val Phe
 370 375 380

Gly Gly Trp Gln Phe Val Arg Phe Val Pro Lys Arg Glu Thr Val Gly
 385 390 395 400

Gly Ala Val Asn Asn Ala Ala Tyr Arg Arg Asn Gly Val Tyr Ala Gly
 405 410 415

Trp Ala Gln Glu Trp Arg Gln Leu Gly Gly Leu Asn Ser Arg Val Ser
 420 425 430

Ala Ser Tyr Ala Arg Arg Asn Tyr Lys Gly Val Ala Ala Phe Ser Thr
 435 440 445

Glu Ala Gln Arg Asn Arg Glu Trp Asn Val Ser Leu Ala Leu Ser His
 450 455 460

Asp Lys Leu Ser Tyr Lys Gly Ile Val Pro Ala Leu Asn Tyr Arg Phe
 465 470 475 480

Gly Arg Thr Glu Ser Asn Val Pro Tyr Ala Lys Arg Arg Asn Ser Glu
 485 490 495

Val Phe Val Ser Ala Asp Trp Arg Phe
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<210> 101

<211> 1476

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1476)

<400> 101

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ggc aaa ctc aaa cgc atc gcg cac ccc gtt tcc ccg cat ttg gaa atg	96
Gly Lys Leu Lys Arg Ile Ala His Pro Val Ser Pro His Leu Glu Met	
20 25 30	
acc gaa atc gcc gac cgc gtg ctg cgc gcc gaa ggg ccg gcg ttg ttg	144
Thr Glu Ile Ala Asp Arg Val Leu Arg Ala Glu Gly Pro Ala Leu Leu	
35 40 45	
ttt gaa cac cca gtt aag ccc gac ggt acg cgc tat gat tat ccc gtg	192
Phe Glu His Pro Val Lys Pro Asp Gly Thr Arg Tyr Asp Tyr Pro Val	
50 55 60	
ttg gca aac ctg ttc ggc acg ccc gaa cgt gtg gcg atg ggc atg ggc	240
Leu Ala Asn Leu Phe Gly Thr Pro Glu Arg Val Ala Met Gly Met Gly	
65 70 75 80	
gcg gac agc gtg tcc aag ctg cgc gaa atc ggg cag acg ctg gcg tat	288
Ala Asp Ser Val Ser Lys Leu Arg Glu Ile Gly Gln Thr Leu Ala Tyr	
85 90 95	
ttg aaa gaa ccc gaa ccg ccc aaa ggc att aaa gac gcg ttt tcc aaa	336
Leu Lys Glu Pro Glu Pro Pro Lys Gly Ile Lys Asp Ala Phe Ser Lys	
100 105 110	
ctg ccg ctc ttg aaa gac att tgg agc atg gcg ccg aac gtg gtg aaa	384
Leu Pro Leu Leu Lys Asp Ile Trp Ser Met Ala Pro Asn Val Val Lys	
115 120 125	
aat gcg ccg tgt cag gaa atc gta tgg gaa ggc gaa gac gtt gat ttg	432
Asn Ala Pro Cys Gln Glu Ile Val Trp Glu Gly Glu Asp Val Asp Leu	
130 135 140	
tat caa ctt ccg att cag cat tgc tgg ccg gaa gac gtt gcg ccg ctg	480
Tyr Gln Leu Pro Ile Gln His Cys Trp Pro Glu Asp Val Ala Pro Leu	
145 150 155 160	
gta acg tgg ggc ttg acc gtc acg cgc ggg ccg cac aaa aaa cgc caa	528
Val Thr Trp Gly Leu Thr Val Thr Arg Gly Pro His Lys Lys Arg Gln	
165 170 175	
aat ctc ggc att tac cgc caa caa tta atc ggc ata aac aag ctg att	576

Asn Leu Gly Ile Tyr Arg Gln Gln Leu Ile Gly Ile Asn Lys Leu Ile	
180 185 190	
atg cgc tgg ctg tgc cat cgc ggc ggc gcg ctg gat tat cag gaa ttc	624
Met Arg Trp Leu Ser His Arg Gly Gly Ala Leu Asp Tyr Gln Glu Phe	
195 200 205	
cgc aaa ctc aat ccc gat acg ccg tat ccc gtc gcc gtc gtg ctc ggc	672
Arg Lys Leu Asn Pro Asp Thr Pro Tyr Pro Val Ala Val Val Leu Gly	
210 215 220	
tgc gac cct gcc acc att ttg ggc gcg gta acg ccc gtt ccc gat acc	720
Cys Asp Pro Ala Thr Ile Leu Gly Ala Val Thr Pro Val Pro Asp Thr	
225 230 235 240	
ttg agc gaa tac cag ttt gcc gga ctg ctg cgc ggt tgc cgg acg gaa	768
Leu Ser Glu Tyr Gln Phe Ala Gly Leu Leu Arg Gly Ser Arg Thr Glu	
245 250 255	
ctg gtg aaa tgt atc ggc aac gat ttg caa gtg cct gcc cgc gca gaa	816
Leu Val Lys Cys Ile Gly Asn Asp Leu Gln Val Pro Ala Arg Ala Glu	
260 265 270	
atc gtg ttg gaa ggc gtc atc cat cca aac gaa acc gcg ttg gaa ggt	864
Ile Val Leu Glu Gly Val Ile His Pro Asn Glu Thr Ala Leu Glu Gly	
275 280 285	
cca tac ggc gac cac acg ggc tat tac aac gag cag gac cat ttc ccc	912
Pro Tyr Gly Asp His Thr Gly Tyr Tyr Asn Glu Gln Asp His Phe Pro	
290 295 300	
gtg ttt acg gtc gaa cgc atc acc atg cgc gaa aac ccg att tac cac	960
Val Phe Thr Val Glu Arg Ile Thr Met Arg Glu Asn Pro Ile Tyr His	
305 310 315 320	
tct acc tac aca ggc aaa ccg ccc gat gaa ccc gcc gtt ttg ggc gtg	1008
Ser Thr Tyr Thr Gly Lys Pro Pro Asp Glu Pro Ala Val Leu Gly Val	
325 330 335	
gcg ttg aac gaa gtg ttc gta ccg ctt ttg caa aag cag ttc ccc gaa	1056
Ala Leu Asn Glu Val Phe Val Pro Leu Leu Gln Lys Gln Phe Pro Glu	
340 345 350	
atc acc gat ttt tac ctg ccg ccc gaa ggc tgc tcc tac cgc atg gcg	1104
Ile Thr Asp Phe Tyr Leu Pro Pro Glu Gly Cys Ser Tyr Arg Met Ala	
355 360 365	
gtg gtg agc atg aaa aaa cag tac gcc gga cac gcc aag cgc gtg atg	1152

Val Val Ser Met Lys Lys Gln Tyr Ala Gly His Ala Lys Arg Val Met
 370 375 380
 atg ggc tgc tgg tcg ttc ctg cgc cag ttt atg tac acc aaa ttc atc 1200
 Met Gly Cys Trp Ser Phe Leu Arg Gln Phe Met Tyr Thr Lys Phe Ile
 385 390 395 400
 att gtg gtg gat gac gat gtg gat gtg cgc gac tgg aaa gaa gtc atc 1248
 Ile Val Val Asp Asp Asp Val Asp Val Arg Asp Trp Lys Glu Val Ile
 405 410 415
 tgg gcg gta acc acg cgc atg gac ccc gtg cgc gat acc gtt ttg atg 1296
 Trp Ala Val Thr Thr Arg Met Asp Pro Val Arg Asp Thr Val Leu Met
 420 425 430
 gaa aac acg ccc atc gac tac ctc gac ttc gcc agc ccc gtc agc gga 1344
 Glu Asn Thr Pro Ile Asp Tyr Leu Asp Phe Ala Ser Pro Val Ser Gly
 435 440 445
 ctt ggc ggc aaa atg ggt ttg gat gcg acc aac aag tgg ccg ggc gaa 1392
 Leu Gly Gly Lys Met Gly Leu Asp Ala Thr Asn Lys Trp Pro Gly Glu
 450 455 460
 acc gac cgc gaa tgg gga cgg gtg att aaa aaa gac cct gcg gtt acg 1440
 Thr Asp Arg Glu Trp Gly Arg Val Ile Lys Lys Asp Pro Ala Val Thr
 465 470 475 480
 gct aag att gat gag att tgg gag gaa ttg ggg ttg 1476
 Ala Lys Ile Asp Glu Ile Trp Glu Glu Leu Gly Leu
 485 490

<210> 102

<211> 492

<212> PRT

<213> Neisseria meningitidis

<400> 102

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 20 25 30
 Thr Glu Ile Ala Asp Arg Val Leu Arg Ala Glu Gly Pro Ala Leu Leu
 35 40 45
 Phe Glu His Pro Val Lys Pro Asp Gly Thr Arg Tyr Asp Tyr Pro Val

50	55	60
Leu Ala Asn Leu Phe Gly Thr Pro Glu Arg Val Ala Met Gly Met Gly		
65	70	75 80
Ala Asp Ser Val Ser Lys Leu Arg Glu Ile Gly Gln Thr Leu Ala Tyr		
	85	90 95
Leu Lys Glu Pro Glu Pro Pro Lys Gly Ile Lys Asp Ala Phe Ser Lys		
	100	105 110
Leu Pro Leu Leu Lys Asp Ile Trp Ser Met Ala Pro Asn Val Val Lys		
	115	120 125
Asn Ala Pro Cys Gln Glu Ile Val Trp Glu Gly Glu Asp Val Asp Leu		
	130	135 140
Tyr Gln Leu Pro Ile Gln His Cys Trp Pro Glu Asp Val Ala Pro Leu		
	145	150 155 160
Val Thr Trp Gly Leu Thr Val Thr Arg Gly Pro His Lys Lys Arg Gln		
	165	170 175
Asn Leu Gly Ile Tyr Arg Gln Gln Leu Ile Gly Ile Asn Lys Leu Ile		
	180	185 190
Met Arg Trp Leu Ser His Arg Gly Gly Ala Leu Asp Tyr Gln Glu Phe		
	195	200 205
Arg Lys Leu Asn Pro Asp Thr Pro Tyr Pro Val Ala Val Val Leu Gly		
	210	215 220
Cys Asp Pro Ala Thr Ile Leu Gly Ala Val Thr Pro Val Pro Asp Thr		
	225	230 235 240
Leu Ser Glu Tyr Gln Phe Ala Gly Leu Leu Arg Gly Ser Arg Thr Glu		
	245	250 255
Leu Val Lys Cys Ile Gly Asn Asp Leu Gln Val Pro Ala Arg Ala Glu		
	260	265 270
Ile Val Leu Glu Gly Val Ile His Pro Asn Glu Thr Ala Leu Glu Gly		
	275	280 285
Pro Tyr Gly Asp His Thr Gly Tyr Tyr Asn Glu Gln Asp His Phe Pro		
	290	295 300
Val Phe Thr Val Glu Arg Ile Thr Met Arg Glu Asn Pro Ile Tyr His		

305 310 315 320
 Ser Thr Tyr Thr Gly Lys Pro Pro Asp Glu Pro Ala Val Leu Gly Val
 325 330 335
 Ala Leu Asn Glu Val Phe Val Pro Leu Leu Gln Lys Gln Phe Pro Glu
 340 345 350
 Ile Thr Asp Phe Tyr Leu Pro Pro Glu Gly Cys Ser Tyr Arg Met Ala
 355 360 365
 Val Val Ser Met Lys Lys Gln Tyr Ala Gly His Ala Lys Arg Val Met
 370 375 380
 Met Gly Cys Trp Ser Phe Leu Arg Gln Phe Met Tyr Thr Lys Phe Ile
 385 390 395 400
 Ile Val Val Asp Asp Asp Val Asp Val Arg Asp Trp Lys Glu Val Ile
 405 410 415
 Trp Ala Val Thr Thr Arg Met Asp Pro Val Arg Asp Thr Val Leu Met
 420 425 430
 Glu Asn Thr Pro Ile Asp Tyr Leu Asp Phe Ala Ser Pro Val Ser Gly
 435 440 445
 Leu Gly Gly Lys Met Gly Leu Asp Ala Thr Asn Lys Trp Pro Gly Glu
 450 455 460
 Thr Asp Arg Glu Trp Gly Arg Val Ile Lys Lys Asp Pro Ala Val Thr
 465 470 475 480
 Ala Lys Ile Asp Glu Ile Trp Glu Glu Leu Gly Leu
 485 490

<210> 103

<211> 1089

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1089)

<400> 103

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1				5					10						15	
acc ctt ttt aac gcg ctg acc caa tcg ggt atc gaa gcg gca aac tat																96
Thr	Leu	Phe	Asn	Ala	Leu	Thr	Gln	Ser	Gly	Ile	Glu	Ala	Ala	Asn	Tyr	
			20					25					30			
cct ttc tgt acc atc gaa ccc aac gtc ggc atc gtc gaa gtc ccc gat																144
Pro	Phe	Cys	Thr	Ile	Glu	Pro	Asn	Val	Gly	Ile	Val	Glu	Val	Pro	Asp	
		35					40					45				
ccg cgt atg gcc gaa ttg gca aaa atc gtc aat ccg caa aaa atg cag																192
Pro	Arg	Met	Ala	Glu	Leu	Ala	Lys	Ile	Val	Asn	Pro	Gln	Lys	Met	Gln	
		50				55					60					
cct gcc atc gtc gaa ttt gtc gat att gcc ggt ttg gtt gca ggc gcg																240
Pro	Ala	Ile	Val	Glu	Phe	Val	Asp	Ile	Ala	Gly	Leu	Val	Ala	Gly	Ala	
	65					70				75				80		
agc aaa ggc gag ggc ttg ggc aac cag ttc ctt gcc aac atc cgc gaa																288
Ser	Lys	Gly	Glu	Gly	Leu	Gly	Asn	Gln	Phe	Leu	Ala	Asn	Ile	Arg	Glu	
			85				90						95			
acc gat gcg att gtg aat gtc gtg cgc tgc ttt gac gac gac aac atc																336
Thr	Asp	Ala	Ile	Val	Asn	Val	Val	Arg	Cys	Phe	Asp	Asp	Asp	Asn	Ile	
		100					105					110				
gtc cac gtt gca ggc cgc gtc gat ccg att gcc gac att gaa acc atc																384
Val	His	Val	Ala	Gly	Arg	Val	Asp	Pro	Ile	Ala	Asp	Ile	Glu	Thr	Ile	
		115				120					125					
ggc aca gag ttg gca ctt gcc gac ctg gca agt gtc gaa aaa gcc atc																432
Gly	Thr	Glu	Leu	Ala	Leu	Ala	Asp	Leu	Ala	Ser	Val	Glu	Lys	Ala	Ile	
	130					135				140						
gtc cgc gaa gaa aaa cgc gcc cgc tca ggc gac aaa gac gcg caa aag																480
Val	Arg	Glu	Glu	Lys	Arg	Ala	Arg	Ser	Gly	Asp	Lys	Asp	Ala	Gln	Lys	
	145				150				155				160			
ctg gtc gat ttg tgc aaa aaa ctg ctg ccg cat ctg gac gaa ggc aaa																528
Leu	Val	Asp	Leu	Cys	Lys	Lys	Leu	Leu	Pro	His	Leu	Asp	Glu	Gly	Lys	
			165				170					175				
ccc gtg cgt tcc ttc ggt ttg gac gcg gaa gaa cgc gcg atg ctc aaa																576
Pro	Val	Arg	Ser	Phe	Gly	Leu	Asp	Ala	Glu	Glu	Arg	Ala	Met	Leu	Lys	
		180					185					190				
ccg ctg ttc ctg ctg acc gcc aaa ccg gcg atg tat gtg ggc aac gtc																624

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Pro Leu Phe Leu Leu Thr Ala Lys Pro Ala Met Tyr Val Gly Asn Val
      195                      200                      205

gcc gaa gac ggt ttt gaa aac aat ccg cac ctc gac cgc ctg aaa gaa 672
Ala Glu Asp Gly Phe Glu Asn Asn Pro His Leu Asp Arg Leu Lys Glu
      210                      215                      220

ttg gcg gca aaa gaa aac gcc ccc gtc gtc gcc gtt tgc gcc gcg atg 720
Leu Ala Ala Lys Glu Asn Ala Pro Val Val Ala Val Cys Ala Ala Met
      225                      230                      235                      240

gag agc gaa att gcc gaa ttg gaa gac gac gaa aaa gcc gag ttc ctc 768
Glu Ser Glu Ile Ala Glu Leu Glu Asp Asp Glu Lys Ala Glu Phe Leu
      245                      250                      255

gcc gaa atg gcc ttg gaa gaa ccg ggc ctg aac cgc ctg att cgt gcc 816
Ala Glu Met Gly Leu Glu Glu Pro Gly Leu Asn Arg Leu Ile Arg Ala
      260                      265                      270

ggt tac gac ctc ttg ggg ctg caa acc tat ttc acc gcc ggt gtg aaa 864
Gly Tyr Asp Leu Leu Gly Leu Gln Thr Tyr Phe Thr Ala Gly Val Lys
      275                      280                      285

gaa gtc cgc gcg tgg acg ata cac aaa ggc gat acc gcc ccg caa gcc 912
Glu Val Arg Ala Trp Thr Ile His Lys Gly Asp Thr Ala Pro Gln Ala
      290                      295                      300

gcc ggc gtg att cac acc gac ttt gag cgc ggc ttc atc cgc gcc caa 960
Ala Gly Val Ile His Thr Asp Phe Glu Arg Gly Phe Ile Arg Ala Gln
      305                      310                      315                      320

gtc att tct tac gac gac ttt gtc tcg ctc ggc ggc gaa gcc aaa gcc 1008
Val Ile Ser Tyr Asp Asp Phe Val Ser Leu Gly Gly Glu Ala Lys Ala
      325                      330                      335

aaa gaa gcc ggc aaa atg cgt gtg gaa ggt aag gaa tac gtc gtg cag 1056
Lys Glu Ala Gly Lys Met Arg Val Glu Gly Lys Glu Tyr Val Val Gln
      340                      345                      350

gac ggc gat gta atg cac ttt ttg ttt aac gtg 1089
Asp Gly Asp Val Met His Phe Leu Phe Asn Val
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<210> 104

<211> 363

<212> PRT

<213> Neisseria meningitidis

<400> 104

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Pro Phe Cys Thr Ile Glu Pro Asn Val Gly Ile Val Glu Val Pro Asp
 35 40 45

Pro Arg Met Ala Glu Leu Ala Lys Ile Val Asn Pro Gln Lys Met Gln
 50 55 60

Pro Ala Ile Val Glu Phe Val Asp Ile Ala Gly Leu Val Ala Gly Ala
 65 70 75 80

Ser Lys Gly Glu Gly Leu Gly Asn Gln Phe Leu Ala Asn Ile Arg Glu
 85 90 95

Thr Asp Ala Ile Val Asn Val Val Arg Cys Phe Asp Asp Asp Asn Ile
 100 105 110

Val His Val Ala Gly Arg Val Asp Pro Ile Ala Asp Ile Glu Thr Ile
 115 120 125

Gly Thr Glu Leu Ala Leu Ala Asp Leu Ala Ser Val Glu Lys Ala Ile
 130 135 140

Val Arg Glu Glu Lys Arg Ala Arg Ser Gly Asp Lys Asp Ala Gln Lys
 145 150 155 160

Leu Val Asp Leu Cys Lys Lys Leu Leu Pro His Leu Asp Glu Gly Lys
 165 170 175

Pro Val Arg Ser Phe Gly Leu Asp Ala Glu Glu Arg Ala Met Leu Lys
 180 185 190

Pro Leu Phe Leu Leu Thr Ala Lys Pro Ala Met Tyr Val Gly Asn Val
 195 200 205

Ala Glu Asp Gly Phe Glu Asn Asn Pro His Leu Asp Arg Leu Lys Glu
 210 215 220

Leu Ala Ala Lys Glu Asn Ala Pro Val Val Ala Val Cys Ala Ala Met
 225 230 235 240

Glu Ser Glu Ile Ala Glu Leu Glu Asp Asp Glu Lys Ala Glu Phe Leu

245 250 255
 Ala Glu Met Gly Leu Glu Glu Pro Gly Leu Asn Arg Leu Ile Arg Ala
 260 265 270
 Gly Tyr Asp Leu Leu Gly Leu Gln Thr Tyr Phe Thr Ala Gly Val Lys
 275 280 285
 Glu Val Arg Ala Trp Thr Ile His Lys Gly Asp Thr Ala Pro Gln Ala
 290 295 300
 Ala Gly Val Ile His Thr Asp Phe Glu Arg Gly Phe Ile Arg Ala Gln
 305 310 315 320
 Val Ile Ser Tyr Asp Asp Phe Val Ser Leu Gly Gly Glu Ala Lys Ala
 325 330 335
 Lys Glu Ala Gly Lys Met Arg Val Glu Gly Lys Glu Tyr Val Val Gln
 340 345 350
 Asp Gly Asp Val Met His Phe Leu Phe Asn Val
 355 360

<210> 105
 <211> 1177
 <212> DNA
 <213> Neisseria meningitidis

<220>
 <221> CDS
 <222> (1)..(1176)

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 gcg gtt tgg ggc gga tgg tct tat ctg aag ccc gag ccg cag gct gct 96
 Ala Val Trp Gly Gly Trp Ser Tyr Leu Lys Pro Glu Pro Gln Ala Ala
 20 25 30

 tat att acg gaa acg gtc agg cgc ggc gac atc agc cgg acg gtt tct 144
 Tyr Ile Thr Glu Thr Val Arg Arg Gly Asp Ile Ser Arg Thr Val Ser
 35 40 45

 gca aca ggg gag att tcg ccg tcc aac ctg gta tcg gtc ggc gcg cag 192

Ala Thr Gly Glu Ile Ser Pro Ser Asn Leu Val Ser Val Gly Ala Gln	
50 55 60	
gca tcg ggg cag att aag aaa ctt tat gtc aaa ctc ggg caa cag gtt	240
Ala Ser Gly Gln Ile Lys Lys Leu Tyr Val Lys Leu Gly Gln Gln Val	
65 70 75 80	
aaa aag ggc gat ttg att gcg gaa atc aat tcg acc tcg cag acc aat	288
Lys Lys Gly Asp Leu Ile Ala Glu Ile Asn Ser Thr Ser Gln Thr Asn	
85 90 95	
acg ctc aat acg gaa aaa tcc aaa ttg gaa acg tat cag gcg aag ctg	336
Thr Leu Asn Thr Glu Lys Ser Lys Leu Glu Thr Tyr Gln Ala Lys Leu	
100 105 110	
gtg tcg gca cag att gca ttg ggc agc gcg gag aag aaa tat aag cgt	384
Val Ser Ala Gln Ile Ala Leu Gly Ser Ala Glu Lys Lys Tyr Lys Arg	
115 120 125	
cag gcg gcg ttg tgg aag gat gat gcg acc gct aaa gaa gat ttg gaa	432
Gln Ala Ala Leu Trp Lys Asp Asp Ala Thr Ala Lys Glu Asp Leu Glu	
130 135 140	
agc gca cag gat gcg ctt gcc gcc gcc aaa gcc aat gtt gcc gag ctg	480
Ser Ala Gln Asp Ala Leu Ala Ala Ala Lys Ala Asn Val Ala Glu Leu	
145 150 155 160	
aag gct cta atc aga cag agc aaa att tcc atc aat acc gcc gag tcg	528
Lys Ala Leu Ile Arg Gln Ser Lys Ile Ser Ile Asn Thr Ala Glu Ser	
165 170 175	
gaa ttg ggc tac acg cgc att acc gca acg atg gac ggc acg gtg gtg	576
Glu Leu Gly Tyr Thr Arg Ile Thr Ala Thr Met Asp Gly Thr Val Val	
180 185 190	
gcg att ctc gtg gaa gag ggg cag act gtg aac gcg gcg cag tct acg	624
Ala Ile Leu Val Glu Glu Gly Gln Thr Val Asn Ala Ala Gln Ser Thr	
195 200 205	
ccg acg att gtc caa ttg gcg aat ctg gat atg atg ttg aac aaa atg	672
Pro Thr Ile Val Gln Leu Ala Asn Leu Asp Met Met Leu Asn Lys Met	
210 215 220	
cag att gcc gag ggc gat att acc aag gtg aag gcg ggg cag gat att	720
Gln Ile Ala Glu Gly Asp Ile Thr Lys Val Lys Ala Gly Gln Asp Ile	
225 230 235 240	
tcg ttt acg att ttg tcc gaa ccg gat acg ccg att aag gcg aag ctc	768

Ser Phe Thr Ile Leu Ser Glu Pro Asp Thr Pro Ile Lys Ala Lys Leu
 245 250 255
 gac agc gtc gac ccc ggg ctg acc acg atg tcg tcg ggc ggc tac aac 816
 Asp Ser Val Asp Pro Gly Leu Thr Thr Met Ser Ser Gly Gly Tyr Asn
 260 265 270
 agc agt acg gat acg gct tcc aat gcg gtc tac tat tat gcc cgt tcg 864
 Ser Ser Thr Asp Thr Ala Ser Asn Ala Val Tyr Tyr Tyr Ala Arg Ser
 275 280 285
 ttt gtg ccg aat ccg gac ggc aaa ctc gcc acg ggg atg acg acg cag 912
 Phe Val Pro Asn Pro Asp Gly Lys Leu Ala Thr Gly Met Thr Thr Gln
 290 295 300
 aat acg gtt gaa atc gac ggt gtg aaa aat gtg ctg att att ccg tcg 960
 Asn Thr Val Glu Ile Asp Gly Val Lys Asn Val Leu Ile Ile Pro Ser
 305 310 315 320
 ctg acc gtg aaa aat cgc ggc ggc agg gcg ttt gtg cgc gtg ttg ggt 1008
 Leu Thr Val Lys Asn Arg Gly Gly Arg Ala Phe Val Arg Val Leu Gly
 325 330 335
 gca gac ggc aag gcg gcg gaa cgc gaa atc ccg acc ggt atg aga gac 1056
 Ala Asp Gly Lys Ala Ala Glu Arg Glu Ile Arg Thr Gly Met Arg Asp
 340 345 350
 agt atg aat acc gaa gta aaa agc ggg ttg aaa gag ggg gac aaa gtg 1104
 Ser Met Asn Thr Glu Val Lys Ser Gly Leu Lys Glu Gly Asp Lys Val
 355 360 365
 gtc atc tcc gaa ata acc gcc gcc gag cag cag gaa agc ggc gaa cgc 1152
 Val Ile Ser Glu Ile Thr Ala Ala Glu Gln Gln Glu Ser Gly Glu Arg
 370 375 380
 gcc cta ggc ggc ccg ccg cgc cga t 1177
 Ala Leu Gly Gly Pro Pro Arg Arg
 385 390

<210> 106

<211> 392

<212> PRT

<213> Neisseria meningitidis

<400> 106

Met Ala Lys Met Met Lys Trp Ala Ala Val Ala Ala Val Ala Ala Ala
 1 5 10 15

Ala Val Trp Gly Gly Trp Ser Tyr Leu Lys Pro Glu Pro Gln Ala Ala
 20 25 30

Tyr Ile Thr Glu Thr Val Arg Arg Gly Asp Ile Ser Arg Thr Val Ser
 35 40 45

Ala Thr Gly Glu Ile Ser Pro Ser Asn Leu Val Ser Val Gly Ala Gln
 50 55 60

Ala Ser Gly Gln Ile Lys Lys Leu Tyr Val Lys Leu Gly Gln Gln Val
 65 70 75 80

Lys Lys Gly Asp Leu Ile Ala Glu Ile Asn Ser Thr Ser Gln Thr Asn
 85 90 95

Thr Leu Asn Thr Glu Lys Ser Lys Leu Glu Thr Tyr Gln Ala Lys Leu
 100 105 110

Val Ser Ala Gln Ile Ala Leu Gly Ser Ala Glu Lys Lys Tyr Lys Arg
 115 120 125

Gln Ala Ala Leu Trp Lys Asp Asp Ala Thr Ala Lys Glu Asp Leu Glu
 130 135 140

Ser Ala Gln Asp Ala Leu Ala Ala Ala Lys Ala Asn Val Ala Glu Leu
 145 150 155 160

Lys Ala Leu Ile Arg Gln Ser Lys Ile Ser Ile Asn Thr Ala Glu Ser
 165 170 175

Glu Leu Gly Tyr Thr Arg Ile Thr Ala Thr Met Asp Gly Thr Val Val
 180 185 190

Ala Ile Leu Val Glu Glu Gly Gln Thr Val Asn Ala Ala Gln Ser Thr
 195 200 205

Pro Thr Ile Val Gln Leu Ala Asn Leu Asp Met Met Leu Asn Lys Met
 210 215 220

Gln Ile Ala Glu Gly Asp Ile Thr Lys Val Lys Ala Gly Gln Asp Ile
 225 230 235 240

Ser Phe Thr Ile Leu Ser Glu Pro Asp Thr Pro Ile Lys Ala Lys Leu
 245 250 255

Asp Ser Val Asp Pro Gly Leu Thr Thr Met Ser Ser Gly Gly Tyr Asn
 260 265 270

Ser Ser Thr Asp Thr Ala Ser Asn Ala Val Tyr Tyr Tyr Ala Arg Ser
 275 280 285

Phe Val Pro Asn Pro Asp Gly Lys Leu Ala Thr Gly Met Thr Thr Gln
 290 295 300

Asn Thr Val Glu Ile Asp Gly Val Lys Asn Val Leu Ile Ile Pro Ser
 305 310 315 320

Leu Thr Val Lys Asn Arg Gly Gly Arg Ala Phe Val Arg Val Leu Gly
 325 330 335

Ala Asp Gly Lys Ala Ala Glu Arg Glu Ile Arg Thr Gly Met Arg Asp
 340 345 350

Ser Met Asn Thr Glu Val Lys Ser Gly Leu Lys Glu Gly Asp Lys Val
 355 360 365

Val Ile Ser Glu Ile Thr Ala Ala Glu Gln Gln Glu Ser Gly Glu Arg
 370 375 380

Ala Leu Gly Gly Pro Pro Arg Arg
 385 390

<210> 107

<211> 1185

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1185)

<400> 107

atg aca gag gac gaa cgt ttc gcg tgg ctg caa ttg gcg ttt acg ccc 48
 Met Thr Glu Asp Glu Arg Phe Ala Trp Leu Gln Leu Ala Phe Thr Pro
 1 5 10 15

tat atc ggc gcg gaa agt ttc ctg ctg ctg atg cgc cgt ttc ggc agc 96
 Tyr Ile Gly Ala Glu Ser Phe Leu Leu Leu Met Arg Arg Phe Gly Ser
 20 25 30

gcg caa aat gcc ctg tcc gca ccg gcg gaa cag gtg gcg gca ctg ata 144
 Ala Gln Asn Ala Leu Ser Ala Pro Ala Glu Gln Val Ala Ala Leu Ile
 35 40 45

cgg cac aaa cag gcg ctt gag gct tgg cgc aat gcg gaa aaa cgc gct	192
Arg His Lys Gln Ala Leu Glu Ala Trp Arg Asn Ala Glu Lys Arg Ala	
50 55 60	
ctg gcg cgg cag gcg gca gaa gcg gcg ttg gaa tgg gaa atg cgg gac	240
Leu Ala Arg Gln Ala Ala Glu Ala Ala Leu Glu Trp Glu Met Arg Asp	
65 70 75 80	
gga tgc cgc ctg atg ctg ctt cag gat gaa gat ttc ccc gaa atg ctg	288
Gly Cys Arg Leu Met Leu Leu Gln Asp Glu Asp Phe Pro Glu Met Leu	
85 90 95	
acg cag ggg ctg acc gcg ccg ccg gtt ttg ttt ttg cgc ggc aac gtg	336
Thr Gln Gly Leu Thr Ala Pro Pro Val Leu Phe Leu Arg Gly Asn Val	
100 105 110	
cgg ctg ctg cac aag cct tcc gcc gcc atc gtc ggc agc cgc cat gcc	384
Arg Leu Leu His Lys Pro Ser Ala Ala Ile Val Gly Ser Arg His Ala	
115 120 125	
acg ccg cag gcg atg cgg att gcc aaa gat ttc ggc aag tcg ttg ggc	432
Thr Pro Gln Ala Met Arg Ile Ala Lys Asp Phe Gly Lys Ser Leu Gly	
130 135 140	
ggg caa aat atc ccc gtt gtg tcg ggt atg gct tcc ggt atc gat act	480
Gly Gln Asn Ile Pro Val Val Ser Gly Met Ala Ser Gly Ile Asp Thr	
145 150 155 160	
gcc gcc cat cag ggc gca ctg gag gca gaa ggc ggc acc atc gcc gtg	528
Ala Ala His Gln Gly Ala Leu Glu Ala Glu Gly Gly Thr Ile Ala Val	
165 170 175	
tgg ggg acg ggt ata gac cgc att tac ccg ccg tcc aat aaa aac ctt	576
Trp Gly Thr Gly Ile Asp Arg Ile Tyr Pro Pro Ser Asn Lys Asn Leu	
180 185 190	
gcc tat gaa atc gcc gaa aaa gga ttg att gtc agc gag ttt ccc atc	624
Ala Tyr Glu Ile Ala Glu Lys Gly Leu Ile Val Ser Glu Phe Pro Ile	
195 200 205	
ggt acg cgc ccg tat gcc ggc aat ttt ccg cgc cgc aac cgc ctg att	672
Gly Thr Arg Pro Tyr Ala Gly Asn Phe Pro Arg Arg Asn Arg Leu Ile	
210 215 220	
gcc gcc ctg tcg caa gta acg ctg gtg gtt gaa gcc gcc ttg gaa tcc	720
Ala Ala Leu Ser Gln Val Thr Leu Val Val Glu Ala Ala Leu Glu Ser	
225 230 235 240	

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ggt tcg ctg att acc gcc ggc ctg gcg gca gaa atg ggg cgc gaa gtg 768
Gly Ser Leu Ile Thr Ala Gly Leu Ala Ala Glu Met Gly Arg Glu Val
                245                250                255

atg gcg gta ccc ggt tcg ata gac aat ccg cac agc aaa ggc tgc cac 816
Met Ala Val Pro Gly Ser Ile Asp Asn Pro His Ser Lys Gly Cys His
                260                265                270

aaa ctg att aaa gac ggc gca aaa ttg gtg gaa tgc ctg gac gac atc 864
Lys Leu Ile Lys Asp Gly Ala Lys Leu Val Glu Cys Leu Asp Asp Ile
                275                280                285

ctg aac gaa tgc ccg ggg cta ttg caa aat acg ggt gct tca tca tat 912
Leu Asn Glu Cys Pro Gly Leu Leu Gln Asn Thr Gly Ala Ser Ser Tyr
                290                295                300

tct ata aat aag gat acg ccc gac acg gga cgc cgc gcc gtt cag acg 960
Ser Ile Asn Lys Asp Thr Pro Asp Thr Gly Arg Arg Ala Val Gln Thr
305                310                315                320

gca tac gcc ccg ccg cct gcc gca aaa atg ccg tct gaa gcg gcg gca 1008
Ala Tyr Ala Pro Pro Pro Ala Ala Lys Met Pro Ser Glu Ala Ala Ala
                325                330                335

ggc ggc aca gcc gtc ggc ggc ata ctg gat aaa atg ggt ttc gac ccc 1056
Gly Gly Thr Ala Val Gly Gly Ile Leu Asp Lys Met Gly Phe Asp Pro
                340                345                350

atc cat ccc gac gtg ctt gcc gga cag ttg gct atg cct gcc gca gat 1104
Ile His Pro Asp Val Leu Ala Gly Gln Leu Ala Met Pro Ala Ala Asp
                355                360                365

ttg tat gcc gca ctg ttg gaa ttg gaa ttg gac ggc agc gtt gcc gca 1152
Leu Tyr Ala Ala Leu Leu Glu Leu Glu Leu Asp Gly Ser Val Ala Ala
                370                375                380

atg ccc agc ggc aga tac cag cgt atc cga act 1185
Met Pro Ser Gly Arg Tyr Gln Arg Ile Arg Thr
385                390                395

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<210> 108

<211> 395

<212> PRT

<213> Neisseria meningitidis

<400> 108

Met Thr Glu Asp Glu Arg Phe Ala Trp Leu Gln Leu Ala Phe Thr Pro
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 Tyr Ile Gly Ala Glu Ser Phe Leu Leu Leu Met Arg Arg Phe Gly Ser
 20 25 30
 Ala Gln Asn Ala Leu Ser Ala Pro Ala Glu Gln Val Ala Ala Leu Ile
 35 40 45
 Arg His Lys Gln Ala Leu Glu Ala Trp Arg Asn Ala Glu Lys Arg Ala
 50 55 60
 Leu Ala Arg Gln Ala Ala Glu Ala Ala Leu Glu Trp Glu Met Arg Asp
 65 70 75 80
 Gly Cys Arg Leu Met Leu Leu Gln Asp Glu Asp Phe Pro Glu Met Leu
 85 90 95
 Thr Gln Gly Leu Thr Ala Pro Pro Val Leu Phe Leu Arg Gly Asn Val
 100 105 110
 Arg Leu Leu His Lys Pro Ser Ala Ala Ile Val Gly Ser Arg His Ala
 115 120 125
 Thr Pro Gln Ala Met Arg Ile Ala Lys Asp Phe Gly Lys Ser Leu Gly
 130 135 140
 Gly Gln Asn Ile Pro Val Val Ser Gly Met Ala Ser Gly Ile Asp Thr
 145 150 155 160
 Ala Ala His Gln Gly Ala Leu Glu Ala Glu Gly Gly Thr Ile Ala Val
 165 170 175
 Trp Gly Thr Gly Ile Asp Arg Ile Tyr Pro Pro Ser Asn Lys Asn Leu
 180 185 190
 Ala Tyr Glu Ile Ala Glu Lys Gly Leu Ile Val Ser Glu Phe Pro Ile
 195 200 205
 Gly Thr Arg Pro Tyr Ala Gly Asn Phe Pro Arg Arg Asn Arg Leu Ile
 210 215 220
 Ala Ala Leu Ser Gln Val Thr Leu Val Val Glu Ala Ala Leu Glu Ser
 225 230 235 240
 Gly Ser Leu Ile Thr Ala Gly Leu Ala Ala Glu Met Gly Arg Glu Val
 245 250 255

Met Ala Val Pro Gly Ser Ile Asp Asn Pro His Ser Lys Gly Cys His
 260 265 270

Lys Leu Ile Lys Asp Gly Ala Lys Leu Val Glu Cys Leu Asp Asp Ile
 275 280 285

Leu Asn Glu Cys Pro Gly Leu Leu Gln Asn Thr Gly Ala Ser Ser Tyr
 290 295 300

Ser Ile Asn Lys Asp Thr Pro Asp Thr Gly Arg Arg Ala Val Gln Thr
 305 310 315 320

Ala Tyr Ala Pro Pro Pro Ala Ala Lys Met Pro Ser Glu Ala Ala Ala
 325 330 335

Gly Gly Thr Ala Val Gly Gly Ile Leu Asp Lys Met Gly Phe Asp Pro
 340 345 350

Ile His Pro Asp Val Leu Ala Gly Gln Leu Ala Met Pro Ala Ala Asp
 355 360 365

Leu Tyr Ala Ala Leu Leu Glu Leu Glu Leu Asp Gly Ser Val Ala Ala
 370 375 380

Met Pro Ser Gly Arg Tyr Gln Arg Ile Arg Thr
 385 390 395

<210> 109

<211> 1257

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1257)

<400> 109

atg aaa cag acc gtc ctc aaa aat aac ctg caa aac ctg ctt gaa agc 48
 Met Lys Gln Thr Val Leu Lys Asn Asn Leu Gln Asn Leu Leu Glu Ser
 1 5 10 15

gca gaa aat atc ctg ctg ctt caa ggc cct gtc ggc gat ttt ttt ctg 96
 Ala Glu Asn Ile Leu Leu Leu Gln Gly Pro Val Gly Asp Phe Phe Leu
 20 25 30

cgc ctt gcc gac tgg ctg act gca aac ggc aaa acc gta cat aaa ttc 144

Arg Leu Ala Asp Trp Leu Thr Ala Asn Gly Lys Thr Val His Lys Phe	
35 40 45	
aac ttt aat gca ggc gac gac tat ttt tat ccg ccc act caa gcg cat	192
Asn Phe Asn Ala Gly Asp Asp Tyr Phe Tyr Pro Pro Thr Gln Ala His	
50 55 60	
acc gtt gtt ttt aac gac aac tac gat gcc ttt cct gag ttt ttg caa	240
Thr Val Val Phe Asn Asp Asn Tyr Asp Ala Phe Pro Glu Phe Leu Gln	
65 70 75 80	
gaa tac atc act caa cat cac atc cag gcc gtt gtc tgc ttt ggc gac	288
Glu Tyr Ile Thr Gln His His Ile Gln Ala Val Val Cys Phe Gly Asp	
85 90 95	
aca cgc cct tat cac gtc att gca aaa cgc att gca aac gaa aac caa	336
Thr Arg Pro Tyr His Val Ile Ala Lys Arg Ile Ala Asn Glu Asn Gln	
100 105 110	
gcc agt ttc tgg gcg ttt gaa gaa ggc tat ttc cgc ccc tac tac atc	384
Ala Ser Phe Trp Ala Phe Glu Glu Gly Tyr Phe Arg Pro Tyr Tyr Ile	
115 120 125	
acc tta gaa aaa gac ggc gtc aac gca ttt tcc ccg ttg ccg cgc cgt	432
Thr Leu Glu Lys Asp Gly Val Asn Ala Phe Ser Pro Leu Pro Arg Arg	
130 135 140	
gcc gac ttt ttt ctt gaa caa ttc cct aag ctt gcc cag caa gaa tat	480
Ala Asp Phe Phe Leu Glu Gln Phe Pro Lys Leu Ala Gln Gln Glu Tyr	
145 150 155 160	
aaa gcg cca acg ccg gta cac ggc ggt ttt acg ccc atg gca aaa aac	528
Lys Ala Pro Thr Pro Val His Gly Gly Phe Thr Pro Met Ala Lys Asn	
165 170 175	
gct atc cgt tac tat atc gag ttg ttc cgc aat cta cgc aaa tac ccc	576
Ala Ile Arg Tyr Tyr Ile Glu Leu Phe Arg Asn Leu Arg Lys Tyr Pro	
180 185 190	
gac tac atc cac cac cgc gca ccc aat gcc ggc cat tac ctc aaa ccg	624
Asp Tyr Ile His His Arg Ala Pro Asn Ala Gly His Tyr Leu Lys Pro	
195 200 205	
tgg tcg ctc tcc atc ctc aag cgt ttg aac tac tat att gaa gac atc	672
Trp Ser Leu Ser Ile Leu Lys Arg Leu Asn Tyr Tyr Ile Glu Asp Ile	
210 215 220	
caa atc gcc aaa cgt gtg gaa gca ggc aaa tac ggc aag ttt ttt att	720

Gln Ile Ala Lys Arg Val Glu Ala Gly Lys Tyr Gly Lys Phe Phe Ile
 225 230 235 240

gtt ccc ttg cag gta ttc aac gac agc caa gtc cgt atc cat tgc gac 768
 Val Pro Leu Gln Val Phe Asn Asp Ser Gln Val Arg Ile His Cys Asp
 245 250 255

ttt ccc agc gtc cgc agc ttc ctg ctc cat gtt ttg agt tca ttt gcc 816
 Phe Pro Ser Val Arg Ser Phe Leu Leu His Val Leu Ser Ser Phe Ala
 260 265 270

gaa cac gcg cct gcc gat acc aac atc atc atc aag cat cat cct atg 864
 Glu His Ala Pro Ala Asp Thr Asn Ile Ile Ile Lys His His Pro Met
 275 280 285

gac cgc ggc ttt atc gac tac tgg cgc gac att aaa cgc ttt atc aaa 912
 Asp Arg Gly Phe Ile Asp Tyr Trp Arg Asp Ile Lys Arg Phe Ile Lys
 290 295 300

gaa cac ccc gaa ctc aaa ggc cgt gtg att tat gtc cat gat gtc ccc 960
 Glu His Pro Glu Leu Lys Gly Arg Val Ile Tyr Val His Asp Val Pro
 305 310 315 320

ctg ccc gtt ttc ctg cgc cac ggt ctc ggc atg gtt acc atc aac agc 1008
 Leu Pro Val Phe Leu Arg His Gly Leu Gly Met Val Thr Ile Asn Ser
 325 330 335

acc agc ggc ctg tcc gga ctg att cac aat atg cca gtt aag gtt ctc 1056
 Thr Ser Gly Leu Ser Gly Leu Ile His Asn Met Pro Val Lys Val Leu
 340 345 350

ggc cgt gcc tat tat gat att ccc ggc att act gac caa aat acc ttg 1104
 Gly Arg Ala Tyr Tyr Asp Ile Pro Gly Ile Thr Asp Gln Asn Thr Leu
 355 360 365

gca gaa ttt tgg aat cat ccg aca ccg cct gac aaa gag ctg ttc cat 1152
 Ala Glu Phe Trp Asn His Pro Thr Pro Pro Asp Lys Glu Leu Phe His
 370 375 380

gcc tac cga atg tac cac ctc aac gtg acc caa att aac ggc aac ttc 1200
 Ala Tyr Arg Met Tyr His Leu Asn Val Thr Gln Ile Asn Gly Asn Phe
 385 390 395 400

tac agt cag gtc ttt ttt cct aac aaa aac aca tcc gac tct tcc aca 1248
 Tyr Ser Gln Val Phe Phe Pro Asn Lys Asn Thr Ser Asp Ser Ser Thr
 405 410 415

cca aca acc 1257

Pro Thr Thr

<210> 110

<211> 419

<212> PRT

<213> Neisseria meningitidis

<400> 110

Met Lys Gln Thr Val Leu Lys Asn Asn Leu Gln Asn Leu Leu Glu Ser

1 5 10 15

Ala Glu Asn Ile Leu Leu Leu Gln Gly Pro Val Gly Asp Phe Phe Leu

20 25 30

Arg Leu Ala Asp Trp Leu Thr Ala Asn Gly Lys Thr Val His Lys Phe

35 40 45

Asn Phe Asn Ala Gly Asp Asp Tyr Phe Tyr Pro Pro Thr Gln Ala His

50 55 60

Thr Val Val Phe Asn Asp Asn Tyr Asp Ala Phe Pro Glu Phe Leu Gln

65 70 75 80

Glu Tyr Ile Thr Gln His His Ile Gln Ala Val Val Cys Phe Gly Asp

85 90 95

Thr Arg Pro Tyr His Val Ile Ala Lys Arg Ile Ala Asn Glu Asn Gln

100 105 110

Ala Ser Phe Trp Ala Phe Glu Glu Gly Tyr Phe Arg Pro Tyr Tyr Ile

115 120 125

Thr Leu Glu Lys Asp Gly Val Asn Ala Phe Ser Pro Leu Pro Arg Arg

130 135 140

Ala Asp Phe Phe Leu Glu Gln Phe Pro Lys Leu Ala Gln Gln Glu Tyr

145 150 155 160

Lys Ala Pro Thr Pro Val His Gly Gly Phe Thr Pro Met Ala Lys Asn

165 170 175

Ala Ile Arg Tyr Tyr Ile Glu Leu Phe Arg Asn Leu Arg Lys Tyr Pro

180 185 190

Asp Tyr Ile His His Arg Ala Pro Asn Ala Gly His Tyr Leu Lys Pro

195 200 205

Trp Ser Leu Ser Ile Leu Lys Arg Leu Asn Tyr Tyr Ile Glu Asp Ile
 210 215 220
 Gln Ile Ala Lys Arg Val Glu Ala Gly Lys Tyr Gly Lys Phe Phe Ile
 225 230 235 240
 Val Pro Leu Gln Val Phe Asn Asp Ser Gln Val Arg Ile His Cys Asp
 245 250 255
 Phe Pro Ser Val Arg Ser Phe Leu Leu His Val Leu Ser Ser Phe Ala
 260 265 270
 Glu His Ala Pro Ala Asp Thr Asn Ile Ile Ile Lys His His Pro Met
 275 280 285
 Asp Arg Gly Phe Ile Asp Tyr Trp Arg Asp Ile Lys Arg Phe Ile Lys
 290 295 300
 Glu His Pro Glu Leu Lys Gly Arg Val Ile Tyr Val His Asp Val Pro
 305 310 315 320
 Leu Pro Val Phe Leu Arg His Gly Leu Gly Met Val Thr Ile Asn Ser
 325 330 335
 Thr Ser Gly Leu Ser Gly Leu Ile His Asn Met Pro Val Lys Val Leu
 340 345 350
 Gly Arg Ala Tyr Tyr Asp Ile Pro Gly Ile Thr Asp Gln Asn Thr Leu
 355 360 365
 Ala Glu Phe Trp Asn His Pro Thr Pro Pro Asp Lys Glu Leu Phe His
 370 375 380
 Ala Tyr Arg Met Tyr His Leu Asn Val Thr Gln Ile Asn Gly Asn Phe
 385 390 395 400
 Tyr Ser Gln Val Phe Phe Pro Asn Lys Asn Thr Ser Asp Ser Ser Thr
 405 410 415
 Pro Thr Thr

<210> 111

<211> 1674

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1674)

<400> 111

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atg agt ttc aaa acc gat gcc gaa atc gcc caa tcc tcc acc atg cgc      48
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   1             5             10             15

ccg att ggc gaa att gcc gcc aag ctg ggt ttg aac gtt gac aac att      96
Pro Ile Gly Glu Ile Ala Ala Lys Leu Gly Leu Asn Val Asp Asn Ile
           20             25             30

gag cct tac ggt cat tac aaa gcc aaa atc aat cct gcc gaa gcg ttc     144
Glu Pro Tyr Gly His Tyr Lys Ala Lys Ile Asn Pro Ala Glu Ala Phe
          35             40             45

aaa ctg ccg caa aaa cag ggc agg ctg att ttg gtt acc gcc atc aac     192
Lys Leu Pro Gln Lys Gln Gly Arg Leu Ile Leu Val Thr Ala Ile Asn
          50             55             60

ccg act ccg gcg ggc gaa ggt aaa acc acc gta acc atc ggt ttg gcg     240
Pro Thr Pro Ala Gly Glu Gly Lys Thr Thr Val Thr Ile Gly Leu Ala
          65             70             75             80

gac gca ttg cgc cat atc ggc aaa gac tct gtg att gct ttg cgc gag     288
Asp Ala Leu Arg His Ile Gly Lys Asp Ser Val Ile Ala Leu Arg Glu
          85             90             95

cct tct ttg ggt ccg gtg ttc ggc gtg aaa ggc ggc gcg gca ggc ggc     336
Pro Ser Leu Gly Pro Val Phe Gly Val Lys Gly Gly Ala Ala Gly Gly
          100            105            110

ggc tat gcc caa gtt ttg ccg atg gaa gac atc aac ctg cac ttc acc     384
Gly Tyr Ala Gln Val Leu Pro Met Glu Asp Ile Asn Leu His Phe Thr
          115            120            125

gga gat ttt cac gcc atc ggt gcg gca aat aat ctg ctt gcc gcg atg     432
Gly Asp Phe His Ala Ile Gly Ala Ala Asn Asn Leu Leu Ala Ala Met
          130            135            140

ctc gac aac cat atc tac caa ggc aac gag ttg aac atc gac ccc aaa     480
Leu Asp Asn His Ile Tyr Gln Gly Asn Glu Leu Asn Ile Asp Pro Lys
          145            150            155            160

cgc gtg ctg tgg cgg cgc gtg gtc gat atg aac gac cgc cag ttg cgc     528
Arg Val Leu Trp Arg Arg Val Val Asp Met Asn Asp Arg Gln Leu Arg

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165	170	175	
aac atc atc gac ggc atg ggc aag cct gtt gac ggc gtg atg cgt cct			576
Asn Ile Ile Asp Gly Met Gly Lys Pro Val Asp Gly Val Met Arg Pro			
180	185	190	
gac ggt ttc gat att acc gtt gct tcc gaa gtg atg gcg gta ttc tgt			624
Asp Gly Phe Asp Ile Thr Val Ala Ser Glu Val Met Ala Val Phe Cys			
195	200	205	
ctt gcc aaa gac atc agc gat ttg aaa gag cgt ttg ggc aac atc ctt			672
Leu Ala Lys Asp Ile Ser Asp Leu Lys Glu Arg Leu Gly Asn Ile Leu			
210	215	220	
gtc gcc tac gcc aaa gac ggc agc ccc gtt tac gcc aaa gat ttg aaa			720
Val Ala Tyr Ala Lys Asp Gly Ser Pro Val Tyr Ala Lys Asp Leu Lys			
225	230	235	240
gcg aat ggc gcg atg gcg gca ttg ctt aaa gat gcg att aag ccc aac			768
Ala Asn Gly Ala Met Ala Ala Leu Leu Lys Asp Ala Ile Lys Pro Asn			
245	250	255	
ttg gtg caa acc atc gaa ggc acg ccc gcc ttc gta cac ggc ggc ccg			816
Leu Val Gln Thr Ile Glu Gly Thr Pro Ala Phe Val His Gly Gly Pro			
260	265	270	
ttc gcc aac atc gcc cac ggc tgc aac tcc gta acc gca acc cgt ctg			864
Phe Ala Asn Ile Ala His Gly Cys Asn Ser Val Thr Ala Thr Arg Leu			
275	280	285	
gcg aaa cac ctt gcc gat tac gcc gta acc gaa gca ggc ttc ggc gcg			912
Ala Lys His Leu Ala Asp Tyr Ala Val Thr Glu Ala Gly Phe Gly Ala			
290	295	300	
gac ttg ggc gcg gaa aaa ttc tgc gac atc aaa tgc cgc ctt gcc ggt			960
Asp Leu Gly Ala Glu Lys Phe Cys Asp Ile Lys Cys Arg Leu Ala Gly			
305	310	315	320
ttg aaa cct gat gcg gct gtt gtc gtg gcg act gtc cgc gcg ttg aaa			1008
Leu Lys Pro Asp Ala Ala Val Val Val Ala Thr Val Arg Ala Leu Lys			
325	330	335	
tat aac ggc ggc gtg gaa cgc gcc aac ctc ggc gaa gaa aat tta gac			1056
Tyr Asn Gly Gly Val Glu Arg Ala Asn Leu Gly Glu Glu Asn Leu Asp			
340	345	350	
gct ttg gaa aaa ggt ttg ccc aac ctg ctg aaa cac att tcc aac ctg			1104
Ala Leu Glu Lys Gly Leu Pro Asn Leu Leu Lys His Ile Ser Asn Leu			

355	360	365	
aaa aac gta ttc gga ctg ccc gtc gtc gtt gcg ctc aac cgc ttc gtg			1152
Lys Asn Val Phe Gly Leu Pro Val Val Val Ala Leu Asn Arg Phe Val			
370	375	380	
tcc gac tcc gat gcc gag ttg gcg atg att gaa aaa gcc tgt gcc gaa			1200
Ser Asp Ser Asp Ala Glu Leu Ala Met Ile Glu Lys Ala Cys Ala Glu			
385	390	395	400
cac ggc gtt gaa gtt tcc ctg acc gaa gtg tgg ggc aaa ggt ggt gcg			1248
His Gly Val Glu Val Ser Leu Thr Glu Val Trp Gly Lys Gly Gly Ala			
405	410	415	
ggc ggc gcg gat ttg gcg cgc aaa gtc gtc aac gcc att gaa agt caa			1296
Gly Gly Ala Asp Leu Ala Arg Lys Val Val Asn Ala Ile Glu Ser Gln			
420	425	430	
acc aat aac ttc ggt ttc gcc tac gat gtc gag ttg ggc atc aaa gac			1344
Thr Asn Asn Phe Gly Phe Ala Tyr Asp Val Glu Leu Gly Ile Lys Asp			
435	440	445	
aaa atc cgt gcg att gcc caa aaa gtg tac ggc gcg gaa gat gtt gat			1392
Lys Ile Arg Ala Ile Ala Gln Lys Val Tyr Gly Ala Glu Asp Val Asp			
450	455	460	
ttc agc gcg gaa gcg tct gcc gaa atc gct tca ctg gaa aaa ctg ggc			1440
Phe Ser Ala Glu Ala Ser Ala Glu Ile Ala Ser Leu Glu Lys Leu Gly			
465	470	475	480
ttg gac aaa atg ccg atc tgc atg gcg aaa acc caa tac tct ttg agc			1488
Leu Asp Lys Met Pro Ile Cys Met Ala Lys Thr Gln Tyr Ser Leu Ser			
485	490	495	
gac aac gcc aaa ctg ttg ggc tgc ccc gaa gac ttc cgc atc gcc gtg			1536
Asp Asn Ala Lys Leu Leu Gly Cys Pro Glu Asp Phe Arg Ile Ala Val			
500	505	510	
cgc ggc atc acc gtt tcc gca ggc gca ggt ttc atc gtc gcc ctg tgc			1584
Arg Gly Ile Thr Val Ser Ala Gly Ala Gly Phe Ile Val Ala Leu Cys			
515	520	525	
ggc aac atg atg aaa atg ccc ggc ctg ccc aaa gtt ccg gct gcc gag			1632
Gly Asn Met Met Lys Met Pro Gly Leu Pro Lys Val Pro Ala Ala Glu			
530	535	540	
aaa atc gat gtg gac gca gaa ggc gtg att cac ggc ttg ttc			1674
Lys Ile Asp Val Asp Ala Glu Gly Val Ile His Gly Leu Phe			

545

550

555

<210> 112

<211> 558

<212> PRT

<213> Neisseria meningitidis

<400> 112

Met Ser Phe Lys Thr Asp Ala Glu Ile Ala Gln Ser Ser Thr Met Arg

1

5

10

15

Pro Ile Gly Glu Ile Ala Ala Lys Leu Gly Leu Asn Val Asp Asn Ile

20

25

30

Glu Pro Tyr Gly His Tyr Lys Ala Lys Ile Asn Pro Ala Glu Ala Phe

35

40

45

Lys Leu Pro Gln Lys Gln Gly Arg Leu Ile Leu Val Thr Ala Ile Asn

50

55

60

Pro Thr Pro Ala Gly Glu Gly Lys Thr Thr Val Thr Ile Gly Leu Ala

65

70

75

80

Asp Ala Leu Arg His Ile Gly Lys Asp Ser Val Ile Ala Leu Arg Glu

85

90

95

Pro Ser Leu Gly Pro Val Phe Gly Val Lys Gly Gly Ala Ala Gly Gly

100

105

110

Gly Tyr Ala Gln Val Leu Pro Met Glu Asp Ile Asn Leu His Phe Thr

115

120

125

Gly Asp Phe His Ala Ile Gly Ala Ala Asn Asn Leu Leu Ala Ala Met

130

135

140

Leu Asp Asn His Ile Tyr Gln Gly Asn Glu Leu Asn Ile Asp Pro Lys

145

150

155

160

Arg Val Leu Trp Arg Arg Val Val Asp Met Asn Asp Arg Gln Leu Arg

165

170

175

Asn Ile Ile Asp Gly Met Gly Lys Pro Val Asp Gly Val Met Arg Pro

180

185

190

Asp Gly Phe Asp Ile Thr Val Ala Ser Glu Val Met Ala Val Phe Cys

195

200

205

Leu Ala Lys Asp Ile Ser Asp Leu Lys Glu Arg Leu Gly Asn Ile Leu
 210 215 220

Val Ala Tyr Ala Lys Asp Gly Ser Pro Val Tyr Ala Lys Asp Leu Lys
 225 230 235 240

Ala Asn Gly Ala Met Ala Ala Leu Leu Lys Asp Ala Ile Lys Pro Asn
 245 250 255

Leu Val Gln Thr Ile Glu Gly Thr Pro Ala Phe Val His Gly Gly Pro
 260 265 270

Phe Ala Asn Ile Ala His Gly Cys Asn Ser Val Thr Ala Thr Arg Leu
 275 280 285

Ala Lys His Leu Ala Asp Tyr Ala Val Thr Glu Ala Gly Phe Gly Ala
 290 295 300

Asp Leu Gly Ala Glu Lys Phe Cys Asp Ile Lys Cys Arg Leu Ala Gly
 305 310 315 320

Leu Lys Pro Asp Ala Ala Val Val Val Ala Thr Val Arg Ala Leu Lys
 325 330 335

Tyr Asn Gly Gly Val Glu Arg Ala Asn Leu Gly Glu Glu Asn Leu Asp
 340 345 350

Ala Leu Glu Lys Gly Leu Pro Asn Leu Leu Lys His Ile Ser Asn Leu
 355 360 365

Lys Asn Val Phe Gly Leu Pro Val Val Val Ala Leu Asn Arg Phe Val
 370 375 380

Ser Asp Ser Asp Ala Glu Leu Ala Met Ile Glu Lys Ala Cys Ala Glu
 385 390 395 400

His Gly Val Glu Val Ser Leu Thr Glu Val Trp Gly Lys Gly Gly Ala
 405 410 415

Gly Gly Ala Asp Leu Ala Arg Lys Val Val Asn Ala Ile Glu Ser Gln
 420 425 430

Thr Asn Asn Phe Gly Phe Ala Tyr Asp Val Glu Leu Gly Ile Lys Asp
 435 440 445

Lys Ile Arg Ala Ile Ala Gln Lys Val Tyr Gly Ala Glu Asp Val Asp
 450 455 460

Phe Ser Ala Glu Ala Ser Ala Glu Ile Ala Ser Leu Glu Lys Leu Gly
465 470 475 480

Leu Asp Lys Met Pro Ile Cys Met Ala Lys Thr Gln Tyr Ser Leu Ser
485 490 495

Asp Asn Ala Lys Leu Leu Gly Cys Pro Glu Asp Phe Arg Ile Ala Val
500 505 510

Arg Gly Ile Thr Val Ser Ala Gly Ala Gly Phe Ile Val Ala Leu Cys
515 520 525

Gly Asn Met Met Lys Met Pro Gly Leu Pro Lys Val Pro Ala Ala Glu
530 535 540

Lys Ile Asp Val Asp Ala Glu Gly Val Ile His Gly Leu Phe
545 550 555

<210> 113

<211> 475

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(474)

<400> 113

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Met Glu His Leu Phe Glu Glu Trp Leu Pro Asp Leu Pro Ala Asp Val
1 5 10 15

tca gac ggc atc ggc ctg ccg atg agc cgg tta ttg aaa gcc cgg tcg 96
Ser Asp Gly Ile Gly Leu Pro Met Ser Arg Leu Leu Lys Ala Arg Ser
20 25 30

ctg act gcc gca ttg cgc gcc ttg ccg cat ctg ttt tcg gta gaa ctg 144
Leu Thr Ala Ala Leu Arg Ala Leu Pro His Leu Phe Ser Val Glu Leu
35 40 45

ctg aaa ctg ggc gaa ttg gaa acg gaa tgc gga ggg cgg ctg gtg cgc 192
Leu Lys Leu Gly Glu Leu Glu Thr Glu Cys Gly Gly Arg Leu Val Arg
50 55 60

gaa gtt ttg ttg aag ctg gac ggt atc cct gtt gtc gcg gca agg agc 240
Glu Val Leu Leu Lys Leu Asp Gly Ile Pro Val Val Ala Ala Arg Ser

65	70	75	80	
gaa tgc cgt atc ggt tgc gcg ttt tgg caa aac att ttg gac tgc ggc				288
Glu Cys Arg Ile Gly Ser Ala Phe Trp Gln Asn Ile Leu Asp Cys Gly				
	85	90	95	
acg cgt cct ttg ggc gaa cgt ctg ttt caa gcc gat ttg gaa ggg gcg				336
Thr Arg Pro Leu Gly Glu Arg Leu Phe Gln Ala Asp Leu Glu Gly Ala				
	100	105	110	
cgt tgc gcg ttt gag ttt gcc gtt ttc ggc gaa gga tgc gga cgg tac				384
Arg Ser Ala Phe Glu Phe Ala Val Phe Gly Glu Gly Cys Gly Arg Tyr				
	115	120	125	
ttt gcc gct cgg cgt tgc cgg ttt tcc cat cac ggc gag gaa atg ctg				432
Phe Ala Ala Arg Arg Ser Arg Phe Ser His His Gly Glu Glu Met Leu				
	130	135	140	
ctg acc gag tat ttt ctg ccc gaa ctg aaa cgc ttt atc gga t				475
Leu Thr Glu Tyr Phe Leu Pro Glu Leu Lys Arg Phe Ile Gly				
145	150	155		
 <210> 114				
<211> 158				
<212> PRT				
<213> Neisseria meningitidis				
 <400> 114				
Met Glu His Leu Phe Glu Glu Trp Leu Pro Asp Leu Pro Ala Asp Val				
1	5	10	15	
 Ser Asp Gly Ile Gly Leu Pro Met Ser Arg Leu Leu Lys Ala Arg Ser				
	20	25	30	
 Leu Thr Ala Ala Leu Arg Ala Leu Pro His Leu Phe Ser Val Glu Leu				
	35	40	45	
 Leu Lys Leu Gly Glu Leu Glu Thr Glu Cys Gly Gly Arg Leu Val Arg				
	50	55	60	
 Glu Val Leu Leu Lys Leu Asp Gly Ile Pro Val Val Ala Ala Arg Ser				
	65	70	75	80
 Glu Cys Arg Ile Gly Ser Ala Phe Trp Gln Asn Ile Leu Asp Cys Gly				
	85	90	95	
 Thr Arg Pro Leu Gly Glu Arg Leu Phe Gln Ala Asp Leu Glu Gly Ala				

100	105	110
Arg Ser Ala Phe Glu Phe Ala Val Phe Gly Glu Gly Cys Gly Arg Tyr		
115	120	125
Phe Ala Ala Arg Arg Ser Arg Phe Ser His His Gly Glu Glu Met Leu		
130	135	140
Leu Thr Glu Tyr Phe Leu Pro Glu Leu Lys Arg Phe Ile Gly		
145	150	155

<210> 115

<211> 1476

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1476)

<400> 115

atg	aaa	tac	aaa	gac	ctg	cgc	gac	ttc	atc	gcc	atg	ctc	gag	cag	cag	48
Met	Lys	Tyr	Lys	Asp	Leu	Arg	Asp	Phe	Ile	Ala	Met	Leu	Glu	Gln	Gln	
1				5					10				15			

ggc	aaa	ctc	aaa	cgc	atc	gcg	cac	ccc	gtt	tcc	ccg	cat	ttg	gaa	atg	96
Gly	Lys	Leu	Lys	Arg	Ile	Ala	His	Pro	Val	Ser	Pro	His	Leu	Glu	Met	
			20					25					30			

acc	gaa	atc	gcc	gac	cgc	gtg	ctg	cgc	gcc	gaa	ggg	ccg	gcg	ttg	ttg	144
Thr	Glu	Ile	Ala	Asp	Arg	Val	Leu	Arg	Ala	Glu	Gly	Pro	Ala	Leu	Leu	
			35				40					45				

ttt	gaa	cac	cca	gtt	aag	ccc	gac	ggt	acg	cgc	tat	gat	tat	ccc	gtg	192
Phe	Glu	His	Pro	Val	Lys	Pro	Asp	Gly	Thr	Arg	Tyr	Asp	Tyr	Pro	Val	
	50					55					60					

ttg	gca	aac	ctg	ttc	ggc	acg	ccc	gaa	cgt	gtg	gcg	atg	ggc	atg	ggc	240
Leu	Ala	Asn	Leu	Phe	Gly	Thr	Pro	Glu	Arg	Val	Ala	Met	Gly	Met	Gly	
65					70					75				80		

gcg	gac	agc	gtg	tcc	aag	ctg	cgc	gaa	atc	ggg	cag	acg	ctg	gcg	tat	288
Ala	Asp	Ser	Val	Ser	Lys	Leu	Arg	Glu	Ile	Gly	Gln	Thr	Leu	Ala	Tyr	
			85					90					95			

ttg	aaa	gaa	ccc	gaa	ccg	ccc	aaa	ggc	att	aaa	gac	gcg	ttt	tcc	aaa	336
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Leu Lys Glu	Pro Glu Pro Pro Lys Gly Ile Lys Asp Ala Phe Ser Lys	
100	105	110
ctg ccg ctc ttg aaa gac att tgg agc atg gcg ccg aac gtg gtg aaa		384
Leu Pro Leu Leu Lys Asp Ile Trp Ser Met Ala Pro Asn Val Val Lys		
115	120	125
aat gcg ccg tgt cag gaa atc gta tgg gaa ggc gaa gac gtt gat ttg		432
Asn Ala Pro Cys Gln Glu Ile Val Trp Glu Gly Glu Asp Val Asp Leu		
130	135	140
tat caa ctt ccg att cag cat tgc tgg ccg gaa gac gtt gcg ccg ctg		480
Tyr Gln Leu Pro Ile Gln His Cys Trp Pro Glu Asp Val Ala Pro Leu		
145	150	155
gta acg tgg ggc ttg acc gtc acg cgc ggg ccg cac aaa aaa cgc caa		528
Val Thr Trp Gly Leu Thr Val Thr Arg Gly Pro His Lys Lys Arg Gln		
165	170	175
aat ctc ggc att tac cgc caa caa tta atc ggc ata aac aag ctg att		576
Asn Leu Gly Ile Tyr Arg Gln Gln Leu Ile Gly Ile Asn Lys Leu Ile		
180	185	190
atg cgc tgg ctg tcg cat cgc ggc ggc gcg ctg gat tat cag gaa ttc		624
Met Arg Trp Leu Ser His Arg Gly Gly Ala Leu Asp Tyr Gln Glu Phe		
195	200	205
cgc aaa ctc aat ccc gat acg ccg tat ccc gtc gcc gtc gtg ctc ggc		672
Arg Lys Leu Asn Pro Asp Thr Pro Tyr Pro Val Ala Val Val Leu Gly		
210	215	220
tgc gac cct gcc acc att ttg ggc gcg gta acg ccc gtt ccc gat acc		720
Cys Asp Pro Ala Thr Ile Leu Gly Ala Val Thr Pro Val Pro Asp Thr		
225	230	235
ttg agc gaa tac cag ttt gcc gga ctg ctg cgc ggt tcg cgg acg gaa		768
Leu Ser Glu Tyr Gln Phe Ala Gly Leu Leu Arg Gly Ser Arg Thr Glu		
245	250	255
ctg gtg aaa tgt atc ggc aac gat ttg caa gtg cct gcc cgc gca gaa		816
Leu Val Lys Cys Ile Gly Asn Asp Leu Gln Val Pro Ala Arg Ala Glu		
260	265	270
atc gtg ttg gaa ggc gtc atc cat cca aac gaa acc gcg ttg gaa ggt		864
Ile Val Leu Glu Gly Val Ile His Pro Asn Glu Thr Ala Leu Glu Gly		
275	280	285
cca tac ggc gac cac acg ggc tat tac aac gag cag gac cat ttc ccc		912

Pro Tyr Gly Asp His Thr Gly Tyr Tyr Asn Glu Gln Asp His Phe Pro	
290 295 300	
gtg ttt acg gtc gaa cgc atc acc atg cgc gaa aac ccg att tac cac	960
Val Phe Thr Val Glu Arg Ile Thr Met Arg Glu Asn Pro Ile Tyr His	
305 310 315 320	
tct acc tac aca ggc aaa ccg ccc gat gaa ccc gcc gtt ttg ggc gtg	1008
Ser Thr Tyr Thr Gly Lys Pro Pro Asp Glu Pro Ala Val Leu Gly Val	
325 330 335	
gcg ttg aac gaa gtg ttc gta ccg ctt ttg caa aag cag ttc ccc gaa	1056
Ala Leu Asn Glu Val Phe Val Pro Leu Leu Gln Lys Gln Phe Pro Glu	
340 345 350	
atc acc gat ttt tac ctg ccg ccc gaa ggc tgc tcc tac cgc atg gcg	1104
Ile Thr Asp Phe Tyr Leu Pro Pro Glu Gly Cys Ser Tyr Arg Met Ala	
355 360 365	
gtg gtg agc atg aaa aaa cag tac gcc gga cac gcc aag cgc gtg atg	1152
Val Val Ser Met Lys Lys Gln Tyr Ala Gly His Ala Lys Arg Val Met	
370 375 380	
atg gcc tgc tgg tcg ttc ctg cgc cag ttt atg tac acc aaa ttc atc	1200
Met Gly Cys Trp Ser Phe Leu Arg Gln Phe Met Tyr Thr Lys Phe Ile	
385 390 395 400	
att gtg gtg gat gac gat gtg gat gtg cgc gac tgg aaa gaa gtc atc	1248
Ile Val Val Asp Asp Asp Val Asp Val Arg Asp Trp Lys Glu Val Ile	
405 410 415	
tgg gcg gta acc acg cgc atg gac ccc gtg cgc gat acc gtt ttg atg	1296
Trp Ala Val Thr Thr Arg Met Asp Pro Val Arg Asp Thr Val Leu Met	
420 425 430	
gaa aac acg ccc atc gac tac ctc gac ttc gcc agc ccc gtc agc gga	1344
Glu Asn Thr Pro Ile Asp Tyr Leu Asp Phe Ala Ser Pro Val Ser Gly	
435 440 445	
ctt ggc ggc aaa atg ggt ttg gat gcg acc aac aag tgg ccg ggc gaa	1392
Leu Gly Gly Lys Met Gly Leu Asp Ala Thr Asn Lys Trp Pro Gly Glu	
450 455 460	
acc gac cgc gaa tgg gga cgg gtg att aaa aaa gac cct gcg gtt acg	1440
Thr Asp Arg Glu Trp Gly Arg Val Ile Lys Lys Asp Pro Ala Val Thr	
465 470 475 480	
gct aag att gat gag att tgg gag gaa ttg ggg ttg	1476

Ala Lys Ile Asp Glu Ile Trp Glu Glu Leu Gly Leu
 485 490

<210> 116

<211> 492

<212> PRT

<213> Neisseria meningitidis

<400> 116

Met Lys Tyr Lys Asp Leu Arg Asp Phe Ile Ala Met Leu Glu Gln Gln
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Gly Lys Leu Lys Arg Ile Ala His Pro Val Ser Pro His Leu Glu Met
 20 25 30

Thr Glu Ile Ala Asp Arg Val Leu Arg Ala Glu Gly Pro Ala Leu Leu
 35 40 45

Phe Glu His Pro Val Lys Pro Asp Gly Thr Arg Tyr Asp Tyr Pro Val
 50 55 60

Leu Ala Asn Leu Phe Gly Thr Pro Glu Arg Val Ala Met Gly Met Gly
 65 70 75 80

Ala Asp Ser Val Ser Lys Leu Arg Glu Ile Gly Gln Thr Leu Ala Tyr
 85 90 95

Leu Lys Glu Pro Glu Pro Pro Lys Gly Ile Lys Asp Ala Phe Ser Lys
 100 105 110

Leu Pro Leu Leu Lys Asp Ile Trp Ser Met Ala Pro Asn Val Val Lys
 115 120 125

Asn Ala Pro Cys Gln Glu Ile Val Trp Glu Gly Glu Asp Val Asp Leu
 130 135 140

Tyr Gln Leu Pro Ile Gln His Cys Trp Pro Glu Asp Val Ala Pro Leu
 145 150 155 160

Val Thr Trp Gly Leu Thr Val Thr Arg Gly Pro His Lys Lys Arg Gln
 165 170 175

Asn Leu Gly Ile Tyr Arg Gln Gln Leu Ile Gly Ile Asn Lys Leu Ile
 180 185 190

Met Arg Trp Leu Ser His Arg Gly Gly Ala Leu Asp Tyr Gln Glu Phe
 195 200 205

Arg Lys Leu Asn Pro Asp Thr Pro Tyr Pro Val Ala Val Val Leu Gly
 210 215 220
 Cys Asp Pro Ala Thr Ile Leu Gly Ala Val Thr Pro Val Pro Asp Thr
 225 230 235 240
 Leu Ser Glu Tyr Gln Phe Ala Gly Leu Leu Arg Gly Ser Arg Thr Glu
 245 250 255
 Leu Val Lys Cys Ile Gly Asn Asp Leu Gln Val Pro Ala Arg Ala Glu
 260 265 270
 Ile Val Leu Glu Gly Val Ile His Pro Asn Glu Thr Ala Leu Glu Gly
 275 280 285
 Pro Tyr Gly Asp His Thr Gly Tyr Tyr Asn Glu Gln Asp His Phe Pro
 290 295 300
 Val Phe Thr Val Glu Arg Ile Thr Met Arg Glu Asn Pro Ile Tyr His
 305 310 315 320
 Ser Thr Tyr Thr Gly Lys Pro Pro Asp Glu Pro Ala Val Leu Gly Val
 325 330 335
 Ala Leu Asn Glu Val Phe Val Pro Leu Leu Gln Lys Gln Phe Pro Glu
 340 345 350
 Ile Thr Asp Phe Tyr Leu Pro Pro Glu Gly Cys Ser Tyr Arg Met Ala
 355 360 365
 Val Val Ser Met Lys Lys Gln Tyr Ala Gly His Ala Lys Arg Val Met
 370 375 380
 Met Gly Cys Trp Ser Phe Leu Arg Gln Phe Met Tyr Thr Lys Phe Ile
 385 390 395 400
 Ile Val Val Asp Asp Asp Val Asp Val Arg Asp Trp Lys Glu Val Ile
 405 410 415
 Trp Ala Val Thr Thr Arg Met Asp Pro Val Arg Asp Thr Val Leu Met
 420 425 430
 Glu Asn Thr Pro Ile Asp Tyr Leu Asp Phe Ala Ser Pro Val Ser Gly
 435 440 445
 Leu Gly Gly Lys Met Gly Leu Asp Ala Thr Asn Lys Trp Pro Gly Glu
 450 455 460

Thr Asp Arg Glu Trp Gly Arg Val Ile Lys Lys Asp Pro Ala Val Thr
 465 470 475 480

Ala Lys Ile Asp Glu Ile Trp Glu Glu Leu Gly Leu
 485 490

<210> 117

<211> 1515

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1515)

<400> 117

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gtt tct gcc gcc tat ggg gcg gat gcg ccc gcg att ttg gat gac aag 96
 Val Ser Ala Ala Tyr Gly Ala Asp Ala Pro Ala Ile Leu Asp Asp Lys
 20 25 30

gca ttg ttg cag gtg cag cgg tcg gtg tcg gat aag tgg gcg gaa tcg 144
 Ala Leu Leu Gln Val Gln Arg Ser Val Ser Asp Lys Trp Ala Glu Ser
 35 40 45

gat tgg aaa gtt gac aat gat gcc ccg cgc gtg gtt gac ggg gat ttt 192
 Asp Trp Lys Val Asp Asn Asp Ala Pro Arg Val Val Asp Gly Asp Phe
 50 55 60

ttg ttg gcg cat ccg aaa atg ttg gaa cat agt ttg cgc gac gtg ctc 240
 Leu Leu Ala His Pro Lys Met Leu Glu His Ser Leu Arg Asp Val Leu
 65 70 75 80

aac ggc aat cag gcg gat ttg atc gct tcg ttg gcg gat ttg tat gcc 288
 Asn Gly Asn Gln Ala Asp Leu Ile Ala Ser Leu Ala Asp Leu Tyr Ala
 85 90 95

aag ctg ccg gat tat gac gcg gtt ttg tac ggc agg gcg cgg gct ttg 336
 Lys Leu Pro Asp Tyr Asp Ala Val Leu Tyr Gly Arg Ala Arg Ala Leu
 100 105 110

ctg gcg aaa ttg gcg gga agg ccg gcg gag gcg gtg gcg cgg tat cgg 384

Leu Ala Lys Leu Ala Gly Arg Pro Ala Glu Ala Val Ala Arg Tyr Arg	
115 120 125	
gaa ctg cac ggg gaa aat gcg gca gac gag cgg att ttg ctg gat ttg	432
Glu Leu His Gly Glu Asn Ala Ala Asp Glu Arg Ile Leu Leu Asp Leu	
130 135 140	
gcg gcg gcg gag ttt gac gat ttc cgg ctg aag tcg gca gaa agg cat	480
Ala Ala Ala Glu Phe Asp Asp Phe Arg Leu Lys Ser Ala Glu Arg His	
145 150 155 160	
ttt gcc gag gcg gaa aaa ttg gat ttg ccg gcg ccg gtt ttg gaa aat	528
Phe Ala Glu Ala Glu Lys Leu Asp Leu Pro Ala Pro Val Leu Glu Asn	
165 170 175	
gtg ggg cgt ttt cgg aaa aaa gcg gag ggg ctg acg ggc tgg cgt ttt	576
Val Gly Arg Phe Arg Lys Lys Ala Glu Gly Leu Thr Gly Trp Arg Phe	
180 185 190	
tcg ggc ggc atc agt ccg gcg gtc aat aga aat gcc aat aat gcc gcg	624
Ser Gly Gly Ile Ser Pro Ala Val Asn Arg Asn Ala Asn Asn Ala Ala	
195 200 205	
ccg cag tat tgc cgg caa aac gga ggc cgg cag ata tgc agt gtc agc	672
Pro Gln Tyr Cys Arg Gln Asn Gly Gly Arg Gln Ile Cys Ser Val Ser	
210 215 220	
cgg gcg gag cgg gcg gca ggc ttg aat tat gaa atc gag gcg gaa aaa	720
Arg Ala Glu Arg Ala Ala Gly Leu Asn Tyr Glu Ile Glu Ala Glu Lys	
225 230 235 240	
ctg acg gcg ttg gca gat aat cat tat ttg ttg ttc cgt tcc aat atc	768
Leu Thr Ala Leu Ala Asp Asn His Tyr Leu Leu Phe Arg Ser Asn Ile	
245 250 255	
ggc ggc acg agc tat tat ttc agt aaa aaa tca gct tat gac gac ggg	816
Gly Gly Thr Ser Tyr Tyr Phe Ser Lys Lys Ser Ala Tyr Asp Asp Gly	
260 265 270	
ttc ggc aga gcg tat ttg ggt tgg cag tat aaa aat gca cgg cag acg	864
Phe Gly Arg Ala Tyr Leu Gly Trp Gln Tyr Lys Asn Ala Arg Gln Thr	
275 280 285	
gcg ggg att ttg ccg ttt tat cag gtg cag ttg tcg ggc agc gac ggc	912
Ala Gly Ile Leu Pro Phe Tyr Gln Val Gln Leu Ser Gly Ser Asp Gly	
290 295 300	
ttt gat gcg aaa aca aaa cgg gta aac aac cgc cgc ctg ccg ccg tat	960

Phe	Asp	Ala	Lys	Thr	Lys	Arg	Val	Asn	Asn	Arg	Arg	Leu	Pro	Pro	Tyr		
305						310					315					320	
atg	ctg	gcg	cac	gga	gtc	ggc	gtg	cag	ttg	tcc	cat	act	tac	cgc	cca	1008	
Met	Leu	Ala	His	Gly	Val	Gly	Val	Gln	Leu	Ser	His	Thr	Tyr	Arg	Pro		
				325					330					335			
aac	ccg	gga	tgg	caa	ttt	tcg	gtc	gcg	ctg	gaa	cat	tac	cgc	caa	cgc	1056	
Asn	Pro	Gly	Trp	Gln	Phe	Ser	Val	Ala	Leu	Glu	His	Tyr	Arg	Gln	Arg		
			340					345					350				
tac	cgc	gaa	cag	gat	agg	gcg	gaa	tac	aat	aac	ggg	cgg	cag	gac	ggg	1104	
Tyr	Arg	Glu	Gln	Asp	Arg	Ala	Glu	Tyr	Asn	Asn	Gly	Arg	Gln	Asp	Gly		
		355				360						365					
ttt	tat	gtt	tcg	tcg	gca	aaa	cgt	ttg	ggc	gaa	tcg	gca	act	gtg	ttc	1152	
Phe	Tyr	Val	Ser	Ser	Ala	Lys	Arg	Leu	Gly	Glu	Ser	Ala	Thr	Val	Phe		
	370					375					380						
ggc	ggc	tgg	cag	ttt	gtg	cgg	ttt	gtg	ccg	aaa	cgc	gaa	acg	gtg	ggc	1200	
Gly	Gly	Trp	Gln	Phe	Val	Arg	Phe	Val	Pro	Lys	Arg	Glu	Thr	Val	Gly		
385				390						395					400		
ggc	gcg	gtc	aat	aat	gcc	gcc	tac	cgg	cgc	aac	ggg	gtt	tat	gcc	ggc	1248	
Gly	Ala	Val	Asn	Asn	Ala	Ala	Tyr	Arg	Arg	Asn	Gly	Val	Tyr	Ala	Gly		
			405					410						415			
tgg	gcg	cag	gag	tgg	cgg	cag	ttg	ggc	ggg	ttg	aac	agt	cgg	gtt	tcc	1296	
Trp	Ala	Gln	Glu	Trp	Arg	Gln	Leu	Gly	Gly	Leu	Asn	Ser	Arg	Val	Ser		
		420						425					430				
gcg	tct	tat	gcc	cgc	cgc	aac	tat	aag	ggc	gtt	gcg	gct	ttc	tcg	aca	1344	
Ala	Ser	Tyr	Ala	Arg	Arg	Asn	Tyr	Lys	Gly	Val	Ala	Ala	Phe	Ser	Thr		
		435				440					445						
gag	gcg	caa	cgc	aac	cgc	gaa	tgg	aat	gtc	tcg	ctg	gct	ttg	agc	cac	1392	
Glu	Ala	Gln	Arg	Asn	Arg	Glu	Trp	Asn	Val	Ser	Leu	Ala	Leu	Ser	His		
	450					455					460						
gac	aag	ttg	tcg	tac	aaa	ggg	atc	gtg	ccc	gcg	ttg	aat	tat	cgt	ttc	1440	
Asp	Lys	Leu	Ser	Tyr	Lys	Gly	Ile	Val	Pro	Ala	Leu	Asn	Tyr	Arg	Phe		
465					470					475					480		
ggc	agg	acg	gaa	agt	aat	gtg	ccg	tat	gcg	aaa	cgc	cgc	aac	agc	gag	1488	
Gly	Arg	Thr	Glu	Ser	Asn	Val	Pro	Tyr	Ala	Lys	Arg	Arg	Asn	Ser	Glu		
			485						490					495			
gtg	ttt	gtg	tcg	gcg	gat	tgg	cgg	ttt								1515	

Val Phe Val Ser Ala Asp Trp Arg Phe
500 505

<210> 118

<211> 505

<212> PRT

<213> Neisseria meningitidis

<400> 118

Met Leu Tyr Phe Arg Tyr Gly Phe Leu Val Val Trp Cys Ala Ala Gly
1 5 10 15

Val Ser Ala Ala Tyr Gly Ala Asp Ala Pro Ala Ile Leu Asp Asp Lys
20 25 30

Ala Leu Leu Gln Val Gln Arg Ser Val Ser Asp Lys Trp Ala Glu Ser
35 40 45

Asp Trp Lys Val Asp Asn Asp Ala Pro Arg Val Val Asp Gly Asp Phe
50 55 60

Leu Leu Ala His Pro Lys Met Leu Glu His Ser Leu Arg Asp Val Leu
65 70 75 80

Asn Gly Asn Gln Ala Asp Leu Ile Ala Ser Leu Ala Asp Leu Tyr Ala
85 90 95

Lys Leu Pro Asp Tyr Asp Ala Val Leu Tyr Gly Arg Ala Arg Ala Leu
100 105 110

Leu Ala Lys Leu Ala Gly Arg Pro Ala Glu Ala Val Ala Arg Tyr Arg
115 120 125

Glu Leu His Gly Glu Asn Ala Ala Asp Glu Arg Ile Leu Leu Asp Leu
130 135 140

Ala Ala Ala Glu Phe Asp Asp Phe Arg Leu Lys Ser Ala Glu Arg His
145 150 155 160

Phe Ala Glu Ala Glu Lys Leu Asp Leu Pro Ala Pro Val Leu Glu Asn
165 170 175

Val Gly Arg Phe Arg Lys Lys Ala Glu Gly Leu Thr Gly Trp Arg Phe
180 185 190

Ser Gly Gly Ile Ser Pro Ala Val Asn Arg Asn Ala Asn Asn Ala Ala
195 200 205

Pro Gln Tyr Cys Arg Gln Asn Gly Gly Arg Gln Ile Cys Ser Val Ser
 210 215 220

Arg Ala Glu Arg Ala Ala Gly Leu Asn Tyr Glu Ile Glu Ala Glu Lys
 225 230 235 240

Leu Thr Ala Leu Ala Asp Asn His Tyr Leu Leu Phe Arg Ser Asn Ile
 245 250 255

Gly Gly Thr Ser Tyr Tyr Phe Ser Lys Lys Ser Ala Tyr Asp Asp Gly
 260 265 270

Phe Gly Arg Ala Tyr Leu Gly Trp Gln Tyr Lys Asn Ala Arg Gln Thr
 275 280 285

Ala Gly Ile Leu Pro Phe Tyr Gln Val Gln Leu Ser Gly Ser Asp Gly
 290 295 300

Phe Asp Ala Lys Thr Lys Arg Val Asn Asn Arg Arg Leu Pro Pro Tyr
 305 310 315 320

Met Leu Ala His Gly Val Gly Val Gln Leu Ser His Thr Tyr Arg Pro
 325 330 335

Asn Pro Gly Trp Gln Phe Ser Val Ala Leu Glu His Tyr Arg Gln Arg
 340 345 350

Tyr Arg Glu Gln Asp Arg Ala Glu Tyr Asn Asn Gly Arg Gln Asp Gly
 355 360 365

Phe Tyr Val Ser Ser Ala Lys Arg Leu Gly Glu Ser Ala Thr Val Phe
 370 375 380

Gly Gly Trp Gln Phe Val Arg Phe Val Pro Lys Arg Glu Thr Val Gly
 385 390 395 400

Gly Ala Val Asn Asn Ala Ala Tyr Arg Arg Asn Gly Val Tyr Ala Gly
 405 410 415

Trp Ala Gln Glu Trp Arg Gln Leu Gly Gly Leu Asn Ser Arg Val Ser
 420 425 430

Ala Ser Tyr Ala Arg Arg Asn Tyr Lys Gly Val Ala Ala Phe Ser Thr
 435 440 445

Glu Ala Gln Arg Asn Arg Glu Trp Asn Val Ser Leu Ala Leu Ser His
 450 455 460

Asp Lys Leu Ser Tyr Lys Gly Ile Val Pro Ala Leu Asn Tyr Arg Phe
 465 470 475 480

Gly Arg Thr Glu Ser Asn Val Pro Tyr Ala Lys Arg Arg Asn Ser Glu
 485 490 495

Val Phe Val Ser Ala Asp Trp Arg Phe
 500 505

<210> 119

<211> 756

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(756)

<400> 119

atg gaa acg cat tcg ggc tgt ttt ttg cgg aag acg gta atg aaa gac 48
 Met Glu Thr His Ser Gly Cys Phe Leu Arg Lys Thr Val Met Lys Asp
 1 5 10 15

gat gtt ttg aaa cag cag gca cac gcg gcg ata cag aag aaa ctg ggc 96
 Asp Val Leu Lys Gln Gln Ala His Ala Ala Ile Gln Lys Lys Leu Gly
 20 25 30

tac gcg ttc cgc gat att tcg ctt ttg cgg cag gct ttg acg cac agg 144
 Tyr Ala Phe Arg Asp Ile Ser Leu Leu Arg Gln Ala Leu Thr His Arg
 35 40 45

agc cat cat gcg aag cac aac gag cgg ttc gag ttt gtc ggc gat tcg 192
 Ser His His Ala Lys His Asn Glu Arg Phe Glu Phe Val Gly Asp Ser
 50 55 60

att ttg aat tat acg gtg gcg cgg atg ctg ttt gac gcg ttt ccg aag 240
 Ile Leu Asn Tyr Thr Val Ala Arg Met Leu Phe Asp Ala Phe Pro Lys
 65 70 75 80

ttg acc gag ggc gag ttg tcg cgg ttg cgg gca agt ctg gtc aat gag 288
 Leu Thr Glu Gly Glu Leu Ser Arg Leu Arg Ala Ser Leu Val Asn Glu
 85 90 95

ggc gtg ctg gcg gaa atg gcg gcg gaa atg aat gtc ggc gac ggc ctg 336
 Gly Val Leu Ala Glu Met Ala Ala Glu Met Asn Val Gly Asp Gly Leu

100	105	110	
tac ttg ggg gcg ggc gag ttg aag agc ggc ggc ttc aga cgg cct tcg			384
Tyr Leu Gly Ala Gly Glu Leu Lys Ser Gly Gly Phe Arg Arg Pro Ser			
115	120	125	
ata ctg gcg gac gcg atg gag gcg atg ttt gcg gcg gtc agc ttc gat			432
Ile Leu Ala Asp Ala Met Glu Ala Met Phe Ala Ala Val Ser Phe Asp			
130	135	140	
gcg gat ttc aac acg gcg gaa aag gtg gtg cgc cat ttg ttt gca gaa			480
Ala Asp Phe Asn Thr Ala Glu Lys Val Val Arg His Leu Phe Ala Glu			
145	150	155	160
cgc gtc cgg cga gtc gat ttc caa aat cag gca aaa gac ggc aaa act			528
Arg Val Arg Arg Val Asp Phe Gln Asn Gln Ala Lys Asp Gly Lys Thr			
165	170	175	
gct ttg cag gag gcg ttg cag gcg cgc cgt ttc gcc ttg ccg aaa tac			576
Ala Leu Gln Glu Ala Leu Gln Ala Arg Arg Phe Ala Leu Pro Lys Tyr			
180	185	190	
cgt atc gaa gag caa atc ggc cat gcc aac gac agt atg ttt gtc att			624
Arg Ile Glu Glu Gln Ile Gly His Ala Asn Asp Ser Met Phe Val Ile			
195	200	205	
tcc tgc gat ttg ggc gaa ctg ggt ttc gtg tgc cgt gcc aaa ggg acg			672
Ser Cys Asp Leu Gly Glu Leu Gly Phe Val Cys Arg Ala Lys Gly Thr			
210	215	220	
agc cgc aag gcg gcg gag cag gaa gcg gcg aaa gag gct ttg aaa tgg			720
Ser Arg Lys Ala Ala Glu Gln Glu Ala Ala Lys Glu Ala Leu Lys Trp			
225	230	235	240
ctg gaa gag aag ctg ccg ctg aag aag aaa aag aaa			756
Leu Glu Glu Lys Leu Pro Leu Lys Lys Lys Lys Lys			
245	250		

<210> 120

<211> 252

<212> PRT

<213> Neisseria meningitidis

<400> 120

Met Glu Thr His Ser Gly Cys Phe Leu Arg Lys Thr Val Met Lys Asp

1

5

10

15

Asp Val Leu Lys Gln Gln Ala His Ala Ala Ile Gln Lys Lys Leu Gly
 20 25 30
 Tyr Ala Phe Arg Asp Ile Ser Leu Leu Arg Gln Ala Leu Thr His Arg
 35 40 45
 Ser His His Ala Lys His Asn Glu Arg Phe Glu Phe Val Gly Asp Ser
 50 55 60
 Ile Leu Asn Tyr Thr Val Ala Arg Met Leu Phe Asp Ala Phe Pro Lys
 65 70 75 80
 Leu Thr Glu Gly Glu Leu Ser Arg Leu Arg Ala Ser Leu Val Asn Glu
 85 90 95
 Gly Val Leu Ala Glu Met Ala Ala Glu Met Asn Val Gly Asp Gly Leu
 100 105 110
 Tyr Leu Gly Ala Gly Glu Leu Lys Ser Gly Gly Phe Arg Arg Pro Ser
 115 120 125
 Ile Leu Ala Asp Ala Met Glu Ala Met Phe Ala Ala Val Ser Phe Asp
 130 135 140
 Ala Asp Phe Asn Thr Ala Glu Lys Val Val Arg His Leu Phe Ala Glu
 145 150 155 160
 Arg Val Arg Arg Val Asp Phe Gln Asn Gln Ala Lys Asp Gly Lys Thr
 165 170 175
 Ala Leu Gln Glu Ala Leu Gln Ala Arg Arg Phe Ala Leu Pro Lys Tyr
 180 185 190
 Arg Ile Glu Glu Gln Ile Gly His Ala Asn Asp Ser Met Phe Val Ile
 195 200 205
 Ser Cys Asp Leu Gly Glu Leu Gly Phe Val Cys Arg Ala Lys Gly Thr
 210 215 220
 Ser Arg Lys Ala Ala Glu Gln Glu Ala Ala Lys Glu Ala Leu Lys Trp
 225 230 235 240
 Leu Glu Glu Lys Leu Pro Leu Lys Lys Lys Lys Lys
 245 250

<211> 570

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(570)

<400> 121

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atg gac aat cac gcc gaa gca cac tgg caa aac ggc tgg ctt caa agc   48
Met Asp Asn His Ala Glu Ala His Trp Gln Asn Gly Trp Leu Gln Ser
   1             5             10             15

ata cgc cat acc ccg tcg ccc aat ttc agc ccg agg gaa acg ggg gaa   96
Ile Arg His Thr Pro Ser Pro Asn Phe Ser Pro Arg Glu Thr Gly Glu
          20             25             30

acg gtt tcc ctg atc gtg ttg cac aac att tca ctg ccg ccg ttc gaa  144
Thr Val Ser Leu Ile Val Leu His Asn Ile Ser Leu Pro Pro Phe Glu
          35             40             45

tac ggc acg gat gct gtg gaa aag ctg ttt gcc aac cgg ctc gac ccc  192
Tyr Gly Thr Asp Ala Val Glu Lys Leu Phe Ala Asn Arg Leu Asp Pro
          50             55             60

aac gga cat ccg ttc ttc agc ctg ata cac act ttg cgc gta tcc agc  240
Asn Gly His Pro Phe Phe Ser Leu Ile His Thr Leu Arg Val Ser Ser
          65             70             75             80

cat ttc tta atc aaa cgc gac ggc aaa acg gtg cag ttc gta tca tgc  288
His Phe Leu Ile Lys Arg Asp Gly Lys Thr Val Gln Phe Val Ser Cys
          85             90             95

ggc gat atg gcg tac cac gcg ggc gta tcc tcg ttt cgc gga cgg gaa  336
Gly Asp Met Ala Tyr His Ala Gly Val Ser Ser Phe Arg Gly Arg Glu
          100            105            110

aaa tgc aac gca ttt tcc atc ggc atc gaa ttg gaa ggc tgc gat ttc  384
Lys Cys Asn Ala Phe Ser Ile Gly Ile Glu Leu Glu Gly Cys Asp Phe
          115            120            125

gaa ccc ttt acc gaa gcg caa tac cgt tcg ctc gaa aca ttg ttg gaa  432
Glu Pro Phe Thr Glu Ala Gln Tyr Arg Ser Leu Glu Thr Leu Leu Glu
          130            135            140

gca ctc tgc cgc cgc tac ccc gtt acc gca gta acc gga cat cag gac  480
Ala Leu Cys Arg Arg Tyr Pro Val Thr Ala Val Thr Gly His Gln Asp
          145            150            155            160

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atc gcg ccc ggc cgc aaa acc gac ccc ggc cac ttt ttc gac tgg cgg 528
 ile ala pro gly arg lys thr asp pro gly his phe phe asp trp arg
 165 170 175

cgg ata cgg gag aaa ggg ttt ccc gta gac aga aat gcc gtc 570
 arg ile arg glu lys gly phe pro val asp arg asn ala val
 180 185 190

<210> 122

<211> 190

<212> PRT

<213> Neisseria meningitidis

<400> 122

Met asp asn his ala glu ala his trp gln asn gly trp leu gln ser
 1 5 10 15

ile arg his thr pro ser pro asn phe ser pro arg glu thr gly glu
 20 25 30

thr val ser leu ile val leu his asn ile ser leu pro pro phe glu
 35 40 45

tyr gly thr asp ala val glu lys leu phe ala asn arg leu asp pro
 50 55 60

asn gly his pro phe phe ser leu ile his thr leu arg val ser ser
 65 70 75 80

his phe leu ile lys arg asp gly lys thr val gln phe val ser cys
 85 90 95

gly asp met ala tyr his ala gly val ser ser phe arg gly arg glu
 100 105 110

lys cys asn ala phe ser ile gly ile glu leu glu gly cys asp phe
 115 120 125

glu pro phe thr glu ala gln tyr arg ser leu glu thr leu leu glu
 130 135 140

ala leu cys arg arg tyr pro val thr ala val thr gly his gln asp
 145 150 155 160

ile ala pro gly arg lys thr asp pro gly his phe phe asp trp arg
 165 170 175

Arg Ile Arg Glu Lys Gly Phe Pro Val Asp Arg Asn Ala Val
 180 185 190

<210> 123

<211> 2274

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(2274)

<400> 123

atg aca aca tta cat ttc tca ggc ttc ccg cgt gtc ggt gcc ttc cgc 48
 Met Thr Thr Leu His Phe Ser Gly Phe Pro Arg Val Gly Ala Phe Arg
 1 5 10 15

gaa ttg aaa ttc gca caa gaa aaa tac tgg cgc aaa gaa atc agc gag 96
 Glu Leu Lys Phe Ala Gln Glu Lys Tyr Trp Arg Lys Glu Ile Ser Glu
 20 25 30

caa gaa ttg ctg gct gtt gct aaa gac ttg cgc gag aaa aac tgg aaa 144
 Gln Glu Leu Leu Ala Val Ala Lys Asp Leu Arg Glu Lys Asn Trp Lys
 35 40 45

cac cag gcc gct gcc aac gcc gat tac gtt gcc gta ggc gat ttc act 192
 His Gln Ala Ala Ala Asn Ala Asp Tyr Val Ala Val Gly Asp Phe Thr
 50 55 60

ttc tac gac cac atc ctc gac ctg caa gtc gcc acc ggc gcg att ccc 240
 Phe Tyr Asp His Ile Leu Asp Leu Gln Val Ala Thr Gly Ala Ile Pro
 65 70 75 80

gcc cgc ttc ggc ttc gac agc caa aac cta tct ttg gaa caa ttc ttc 288
 Ala Arg Phe Gly Phe Asp Ser Gln Asn Leu Ser Leu Glu Gln Phe Phe
 85 90 95

caa ctg gcg cgc ggt aac aaa gac caa ttc gct atc gaa atg acc aaa 336
 Gln Leu Ala Arg Gly Asn Lys Asp Gln Phe Ala Ile Glu Met Thr Lys
 100 105 110

ttg ttc gac acc aac tac cac tac ttg gtg cct gaa ttc cac gcc gat 384
 Trp Phe Asp Thr Asn Tyr His Tyr Leu Val Pro Glu Phe His Ala Asp
 115 120 125

acc gaa ttt aaa gcc aat gcc aaa cac tac gtt caa cag ctg caa gaa 432
 Thr Glu Phe Lys Ala Asn Ala Lys His Tyr Val Gln Gln Leu Gln Glu
 130 135 140

gcc caa gct ttg ggc ctg aaa gcc aaa ccg acc gtt gtg ggt ccg ttg 480
 Ala Gln Ala Leu Gly Leu Lys Ala Lys Pro Thr Val Val Gly Pro Leu
 145 150 155 160

act ttc ctg tgg gtc ggt aaa gaa aaa ggc gcc gtt gaa ttc gac cgt 528
 Thr Phe Leu Trp Val Gly Lys Glu Lys Gly Ala Val Glu Phe Asp Arg
 165 170 175

ctg agc ctg ttg cct aaa ctg ttg cct gtt tac gtt gaa atc ctg act 576
 Leu Ser Leu Leu Pro Lys Leu Leu Pro Val Tyr Val Glu Ile Leu Thr
 180 185 190

gct ttg gtt gaa gcc ggt gcc gag tgg att caa atc gac gag cct gct 624
 Ala Leu Val Glu Ala Gly Ala Glu Trp Ile Gln Ile Asp Glu Pro Ala
 195 200 205

ttg act gtt gac ctg cct aaa gaa tgg gta gaa gca tac aaa gac gtt 672
 Leu Thr Val Asp Leu Pro Lys Glu Trp Val Glu Ala Tyr Lys Asp Val
 210 215 220

tac gcc act ttg agc aaa gta agt gcc aaa atc ctg ctg agc act tac 720
 Tyr Ala Thr Leu Ser Lys Val Ser Ala Lys Ile Leu Leu Ser Thr Tyr
 225 230 235 240

ttc ggt tct gtt gcc gaa cac gcc gca ttg ttg aaa tcc ctg cct gtt 768
 Phe Gly Ser Val Ala Glu His Ala Ala Leu Leu Lys Ser Leu Pro Val
 245 250 255

gac ggc ctg cac atc gac ttg gta cgc gct cct gaa caa ctg gat gcg 816
 Asp Gly Leu His Ile Asp Leu Val Arg Ala Pro Glu Gln Leu Asp Ala
 260 265 270

ttc gcc gac tac gac aaa gtt ctg tct gcc ggc gtc att gac ggc cgc 864
 Phe Ala Asp Tyr Asp Lys Val Leu Ser Ala Gly Val Ile Asp Gly Arg
 275 280 285

aat att tgg cgc gcc aac ctg aac aaa gtt ttg gaa act gtt gag ctt 912
 Asn Ile Trp Arg Ala Asn Leu Asn Lys Val Leu Glu Thr Val Glu Leu
 290 295 300

ctg caa gcc aaa ctg ggc gac cgt ttg tgg att tcc agc tct tgc tcg 960
 Leu Gln Ala Lys Leu Gly Asp Arg Leu Trp Ile Ser Ser Ser Cys Ser
 305 310 315 320

ctg ctg cac act ccg ttt gac ttg tca gtt gaa gaa aaa ctg aaa gcc	1008
Leu Leu His Thr Pro Phe Asp Leu Ser Val Glu Glu Lys Leu Lys Ala	
325 330 335	
aac aaa ccc gac ctg tac tct tgg ttg gca ttc acc ctg caa aaa acc	1056
Asn Lys Pro Asp Leu Tyr Ser Trp Leu Ala Phe Thr Leu Gln Lys Thr	
340 345 350	
caa gaa ttg cgc gtc ctg aaa gcc gca ttg aac gaa ggc cgt gat tct	1104
Gln Glu Leu Arg Val Leu Lys Ala Ala Leu Asn Glu Gly Arg Asp Ser	
355 360 365	
gtt gcc gaa gaa ctg gct gcc agc caa gct gct gcc gac tcc cgt gcc	1152
Val Ala Glu Glu Leu Ala Ala Ser Gln Ala Ala Ala Asp Ser Arg Ala	
370 375 380	
aac agc agc gaa atc cac cgt gca gac gtt gcc aaa cgc ctg gcc gat	1200
Asn Ser Ser Glu Ile His Arg Ala Asp Val Ala Lys Arg Leu Ala Asp	
385 390 395 400	
ttg cct gcc aac gca gac caa cgc aaa tcc cca ttt gcc gac cgt atc	1248
Leu Pro Ala Asn Ala Asp Gln Arg Lys Ser Pro Phe Ala Asp Arg Ile	
405 410 415	
aaa gcg caa caa gca tgg ttg aac ctg cct ctg ctg ccg acg acc aac	1296
Lys Ala Gln Gln Ala Trp Leu Asn Leu Pro Leu Leu Pro Thr Thr Asn	
420 425 430	
atc ggt tct ttc ccg caa act acc gaa atc cgc caa gca cgc gca gcc	1344
Ile Gly Ser Phe Pro Gln Thr Thr Glu Ile Arg Gln Ala Arg Ala Ala	
435 440 445	
ttc aaa aaa ggc gaa ctg tct gcc gcc gat tac gaa gcc gcg atg aaa	1392
Phe Lys Lys Gly Glu Leu Ser Ala Ala Asp Tyr Glu Ala Ala Met Lys	
450 455 460	
aaa gaa atc gcc ttg gtg gtt gaa gag caa gaa aaa ctg gac ttg gac	1440
Lys Glu Ile Ala Leu Val Val Glu Glu Gln Glu Lys Leu Asp Leu Asp	
465 470 475 480	
gta ctg gta cac ggc gaa gcc gag cgt aac gac atg gtt gaa tac ttc	1488
Val Leu Val His Gly Glu Ala Glu Arg Asn Asp Met Val Glu Tyr Phe	
485 490 495	
ggc gaa ttg ttg agc ggt ttt gca ttc acc caa tat ggc tgg gta caa	1536
Gly Glu Leu Leu Ser Gly Phe Ala Phe Thr Gln Tyr Gly Trp Val Gln	
500 505 510	

agc tac ggt tct cgc tgc gtt aaa cca ccg att atc ttc ggt gac gta	1584
Ser Tyr Gly Ser Arg Cys Val Lys Pro Pro Ile Ile Phe Gly Asp Val	
515 520 525	
agc cgt cct gaa gcc atg acc gta gct tgg tct act tac gca caa agc	1632
Ser Arg Pro Glu Ala Met Thr Val Ala Trp Ser Thr Tyr Ala Gln Ser	
530 535 540	
ctg acc aaa cgc ccg atg aaa ggt atg ttg acc ggc cct gta acc att	1680
Leu Thr Lys Arg Pro Met Lys Gly Met Leu Thr Gly Pro Val Thr Ile	
545 550 555 560	
ctg caa tgg tct ttc gtc cgc aac gat att cct cgc tct acc gtg tgt	1728
Leu Gln Trp Ser Phe Val Arg Asn Asp Ile Pro Arg Ser Thr Val Cys	
565 570 575	
aaa caa atc gca ctg gct ctg aac gac gaa gta ttg gat ctg gaa aaa	1776
Lys Gln Ile Ala Leu Ala Leu Asn Asp Glu Val Leu Asp Leu Glu Lys	
580 585 590	
gca ggc atc aaa gtc atc caa atc gac gaa cct gcc atc cgc gaa ggc	1824
Ala Gly Ile Lys Val Ile Gln Ile Asp Glu Pro Ala Ile Arg Glu Gly	
595 600 605	
ctg cct ttg aaa cgt gcc gat tgg gat gcc tac ctg aac tgg gcc ggc	1872
Leu Pro Leu Lys Arg Ala Asp Trp Asp Ala Tyr Leu Asn Trp Ala Gly	
610 615 620	
gaa tct ttc cgc ctg tcc tct acc ggt tgc gaa gac agc act caa atc	1920
Glu Ser Phe Arg Leu Ser Ser Thr Gly Cys Glu Asp Ser Thr Gln Ile	
625 630 635 640	
cac acc cat atg tgc tac tct gag ttc aac gac atc ctg cct gcg att	1968
His Thr His Met Cys Tyr Ser Glu Phe Asn Asp Ile Leu Pro Ala Ile	
645 650 655	
gct gca atg gat gcg gac gtg atc acc atc gag act tca cgt tcc gac	2016
Ala Ala Met Asp Ala Asp Val Ile Thr Ile Glu Thr Ser Arg Ser Asp	
660 665 670	
atg gaa ctg ttg acc gcg ttc ggc gaa ttc aaa tac ccg aac gac atc	2064
Met Glu Leu Leu Thr Ala Phe Gly Glu Phe Lys Tyr Pro Asn Asp Ile	
675 680 685	
ggc ccg ggg gtt tac gac atc cac agc ccg cgc gta ccg aca gaa gcc	2112
Gly Pro Gly Val Tyr Asp Ile His Ser Pro Arg Val Pro Thr Glu Ala	
690 695 700	

gaa gtg gag cac ctg ttg cgc aaa gcc atc gag gtt gta ccg gtt gaa 2160
 Glu Val Glu His Leu Leu Arg Lys Ala Ile Glu Val Val Pro Val Glu
 705 710 715 720

cgt ctg tgg gtt aac ccg gac tgc ggc ctg aaa aca cgc ggc tgg aaa 2208
 Arg Leu Trp Val Asn Pro Asp Cys Gly Leu Lys Thr Arg Gly Trp Lys
 725 730 735

gaa act ctg gaa caa ctc caa gtg atg atg aac gta acc cac aaa ttg 2256
 Glu Thr Leu Glu Gln Leu Gln Val Met Met Asn Val Thr His Lys Leu
 740 745 750

cgt gcc gaa ttg gcg aaa 2274
 Arg Ala Glu Leu Ala Lys
 755

<210> 124

<211> 758

<212> PRT

<213> Neisseria meningitidis

<400> 124

Met Thr Thr Leu His Phe Ser Gly Phe Pro Arg Val Gly Ala Phe Arg
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Glu Leu Lys Phe Ala Gln Glu Lys Tyr Trp Arg Lys Glu Ile Ser Glu
 20 25 30

Gln Glu Leu Leu Ala Val Ala Lys Asp Leu Arg Glu Lys Asn Trp Lys
 35 40 45

His Gln Ala Ala Ala Asn Ala Asp Tyr Val Ala Val Gly Asp Phe Thr
 50 55 60

Phe Tyr Asp His Ile Leu Asp Leu Gln Val Ala Thr Gly Ala Ile Pro
 65 70 75 80

Ala Arg Phe Gly Phe Asp Ser Gln Asn Leu Ser Leu Glu Gln Phe Phe
 85 90 95

Gln Leu Ala Arg Gly Asn Lys Asp Gln Phe Ala Ile Glu Met Thr Lys
 100 105 110

Trp Phe Asp Thr Asn Tyr His Tyr Leu Val Pro Glu Phe His Ala Asp
 115 120 125

Thr Glu Phe Lys Ala Asn Ala Lys His Tyr Val Gln Gln Leu Gln Glu

130	135	140
Ala Gln Ala Leu Gly Leu Lys Ala Lys Pro Thr Val Val Gly Pro Leu		
145	150	155 160
Thr Phe Leu Trp Val Gly Lys Glu Lys Gly Ala Val Glu Phe Asp Arg		
	165	170 175
Leu Ser Leu Leu Pro Lys Leu Leu Pro Val Tyr Val Glu Ile Leu Thr		
	180	185 190
Ala Leu Val Glu Ala Gly Ala Glu Trp Ile Gln Ile Asp Glu Pro Ala		
	195	200 205
Leu Thr Val Asp Leu Pro Lys Glu Trp Val Glu Ala Tyr Lys Asp Val		
210	215	220
Tyr Ala Thr Leu Ser Lys Val Ser Ala Lys Ile Leu Leu Ser Thr Tyr		
225	230	235 240
Phe Gly Ser Val Ala Glu His Ala Ala Leu Leu Lys Ser Leu Pro Val		
	245	250 255
Asp Gly Leu His Ile Asp Leu Val Arg Ala Pro Glu Gln Leu Asp Ala		
	260	265 270
Phe Ala Asp Tyr Asp Lys Val Leu Ser Ala Gly Val Ile Asp Gly Arg		
	275	280 285
Asn Ile Trp Arg Ala Asn Leu Asn Lys Val Leu Glu Thr Val Glu Leu		
290	295	300
Leu Gln Ala Lys Leu Gly Asp Arg Leu Trp Ile Ser Ser Ser Cys Ser		
305	310	315 320
Leu Leu His Thr Pro Phe Asp Leu Ser Val Glu Glu Lys Leu Lys Ala		
	325	330 335
Asn Lys Pro Asp Leu Tyr Ser Trp Leu Ala Phe Thr Leu Gln Lys Thr		
	340	345 350
Gln Glu Leu Arg Val Leu Lys Ala Ala Leu Asn Glu Gly Arg Asp Ser		
	355	360 365
Val Ala Glu Glu Leu Ala Ala Ser Gln Ala Ala Ala Asp Ser Arg Ala		
370	375	380
Asn Ser Ser Glu Ile His Arg Ala Asp Val Ala Lys Arg Leu Ala Asp		

385	390	395	400
Leu Pro Ala Asn Ala Asp Gln Arg Lys Ser Pro Phe Ala Asp Arg Ile	405	410	415
Lys Ala Gln Gln Ala Trp Leu Asn Leu Pro Leu Leu Pro Thr Thr Asn	420	425	430
Ile Gly Ser Phe Pro Gln Thr Thr Glu Ile Arg Gln Ala Arg Ala Ala	435	440	445
Phe Lys Lys Gly Glu Leu Ser Ala Ala Asp Tyr Glu Ala Ala Met Lys	450	455	460
Lys Glu Ile Ala Leu Val Val Glu Glu Gln Glu Lys Leu Asp Leu Asp	465	470	475
Val Leu Val His Gly Glu Ala Glu Arg Asn Asp Met Val Glu Tyr Phe	485	490	495
Gly Glu Leu Leu Ser Gly Phe Ala Phe Thr Gln Tyr Gly Trp Val Gln	500	505	510
Ser Tyr Gly Ser Arg Cys Val Lys Pro Pro Ile Ile Phe Gly Asp Val	515	520	525
Ser Arg Pro Glu Ala Met Thr Val Ala Trp Ser Thr Tyr Ala Gln Ser	530	535	540
Leu Thr Lys Arg Pro Met Lys Gly Met Leu Thr Gly Pro Val Thr Ile	545	550	555
Leu Gln Trp Ser Phe Val Arg Asn Asp Ile Pro Arg Ser Thr Val Cys	565	570	575
Lys Gln Ile Ala Leu Ala Leu Asn Asp Glu Val Leu Asp Leu Glu Lys	580	585	590
Ala Gly Ile Lys Val Ile Gln Ile Asp Glu Pro Ala Ile Arg Glu Gly	595	600	605
Leu Pro Leu Lys Arg Ala Asp Trp Asp Ala Tyr Leu Asn Trp Ala Gly	610	615	620
Glu Ser Phe Arg Leu Ser Ser Thr Gly Cys Glu Asp Ser Thr Gln Ile	625	630	635
His Thr His Met Cys Tyr Ser Glu Phe Asn Asp Ile Leu Pro Ala Ile			

645 650 655
 Ala Ala Met Asp Ala Asp Val Ile Thr Ile Glu Thr Ser Arg Ser Asp
 660 665 670
 Met Glu Leu Leu Thr Ala Phe Gly Glu Phe Lys Tyr Pro Asn Asp Ile
 675 680 685
 Gly Pro Gly Val Tyr Asp Ile His Ser Pro Arg Val Pro Thr Glu Ala
 690 695 700
 Glu Val Glu His Leu Leu Arg Lys Ala Ile Glu Val Val Pro Val Glu
 705 710 715 720
 Arg Leu Trp Val Asn Pro Asp Cys Gly Leu Lys Thr Arg Gly Trp Lys
 725 730 735
 Glu Thr Leu Glu Gln Leu Gln Val Met Met Asn Val Thr His Lys Leu
 740 745 750
 Arg Ala Glu Leu Ala Lys
 755

<210> 125
 <211> 1137
 <212> DNA
 <213> Neisseria meningitidis

<220>
 <221> CDS
 <222> (1)..(1137)

<400> 125
 atg agc cgt tta tgg ttt ttt gcc gca aaa aac att ata atc cgc ctt 48
 Met Ser Arg Leu Trp Phe Phe Ala Ala Lys Asn Ile Ile Ile Arg Leu
 1 5 10 15
 att tac cta ttg ccc aag gag aca caa atg gca ctc gta tcc atg cgc 96
 Ile Tyr Leu Leu Pro Lys Glu Thr Gln Met Ala Leu Val Ser Met Arg
 20 25 30
 caa ctg ctt gat cat gct gcc gaa aac agc tac ggc ctg ccc gcg ttc 144
 Gln Leu Leu Asp His Ala Ala Glu Asn Ser Tyr Gly Leu Pro Ala Phe
 35 40 45
 aac gtc aac aac ctc gaa caa atg cgc gcc att atg gaa gcc gcc gac 192

Asn Val Asn Asn Leu Glu Gln Met Arg Ala Ile Met Glu Ala Ala Asp	
50 55 60	
caa gtc aac gcg ccc gtc atc gta cag gcg agc gca ggt gcg cgc aaa	240
Gln Val Asn Ala Pro Val Ile Val Gln Ala Ser Ala Gly Ala Arg Lys	
65 70 75 80	
tac gcg ggc gcg ccg ttt ttg cgc cac ctg att ttg gcg gct gtc gaa	288
Tyr Ala Gly Ala Pro Phe Leu Arg His Leu Ile Leu Ala Ala Val Glu	
85 90 95	
gaa ttt ccg cac atc ccc gtc gtg atg cac caa gac cac ggc gca tcg	336
Glu Phe Pro His Ile Pro Val Val Met His Gln Asp His Gly Ala Ser	
100 105 110	
ccc gac gtg tgc caa cgc tcc atc caa ctg ggc ttt tcc tcc gtg atg	384
Pro Asp Val Cys Gln Arg Ser Ile Gln Leu Gly Phe Ser Ser Val Met	
115 120 125	
atg gac ggc tcg ctg atg gaa gac ggc aaa acc cct tct tct tat gaa	432
Met Asp Gly Ser Leu Met Glu Asp Gly Lys Thr Pro Ser Ser Tyr Glu	
130 135 140	
tac aac gtc aac gcc acc cgt acc gtg gtt aat ttc tcc cac gcc tgc	480
Tyr Asn Val Asn Ala Thr Arg Thr Val Val Asn Phe Ser His Ala Cys	
145 150 155 160	
ggc gta tcc gtt gaa ggc gaa atc ggc gta ttg ggc aac ctc gaa act	528
Gly Val Ser Val Glu Gly Glu Ile Gly Val Leu Gly Asn Leu Glu Thr	
165 170 175	
ggc gaa gcc ggc gaa gaa gac ggt gta ggc gca gtg ggc aaa ctt tcc	576
Gly Glu Ala Gly Glu Glu Asp Gly Val Gly Ala Val Gly Lys Leu Ser	
180 185 190	
cac gac caa atg ctc acc agc gtc gaa gat gcc gtg cgt ttc gtt aaa	624
His Asp Gln Met Leu Thr Ser Val Glu Asp Ala Val Arg Phe Val Lys	
195 200 205	
gat acc ggc gtt gac gca ttg gcg att gcc gtc ggc acc agc cac ggc	672
Asp Thr Gly Val Asp Ala Leu Ala Ile Ala Val Gly Thr Ser His Gly	
210 215 220	
gcg tac aaa ttc acc cgt ccg ccc aca ggc gac gtg ttg cgt atc gac	720
Ala Tyr Lys Phe Thr Arg Pro Pro Thr Gly Asp Val Leu Arg Ile Asp	
225 230 235 240	
cgc atc aaa gaa atc cac caa gcc ctg ccc aat aca cac atc gtg atg	768

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<400> 126
Met Ser Arg Leu Trp Phe Phe Ala Ala Lys Asn Ile Ile Ile Arg Leu
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Ile Tyr Leu Leu Pro Lys Glu Thr Gln Met Ala Leu Val Ser Met Arg
          20             25             30

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Gln Leu Leu Asp His Ala Ala Glu Asn Ser Tyr Gly Leu Pro Ala Phe
 35 40 45
 Asn Val Asn Asn Leu Glu Gln Met Arg Ala Ile Met Glu Ala Ala Asp
 50 55 60
 Gln Val Asn Ala Pro Val Ile Val Gln Ala Ser Ala Gly Ala Arg Lys
 65 70 75 80
 Tyr Ala Gly Ala Pro Phe Leu Arg His Leu Ile Leu Ala Ala Val Glu
 85 90 95
 Glu Phe Pro His Ile Pro Val Val Met His Gln Asp His Gly Ala Ser
 100 105 110
 Pro Asp Val Cys Gln Arg Ser Ile Gln Leu Gly Phe Ser Ser Val Met
 115 120 125
 Met Asp Gly Ser Leu Met Glu Asp Gly Lys Thr Pro Ser Ser Tyr Glu
 130 135 140
 Tyr Asn Val Asn Ala Thr Arg Thr Val Val Asn Phe Ser His Ala Cys
 145 150 155 160
 Gly Val Ser Val Glu Gly Glu Ile Gly Val Leu Gly Asn Leu Glu Thr
 165 170 175
 Gly Glu Ala Gly Glu Glu Asp Gly Val Gly Ala Val Gly Lys Leu Ser
 180 185 190
 His Asp Gln Met Leu Thr Ser Val Glu Asp Ala Val Arg Phe Val Lys
 195 200 205
 Asp Thr Gly Val Asp Ala Leu Ala Ile Ala Val Gly Thr Ser His Gly
 210 215 220
 Ala Tyr Lys Phe Thr Arg Pro Pro Thr Gly Asp Val Leu Arg Ile Asp
 225 230 235 240
 Arg Ile Lys Glu Ile His Gln Ala Leu Pro Asn Thr His Ile Val Met
 245 250 255
 His Gly Ser Ser Ser Val Pro Gln Glu Trp Leu Lys Val Ile Asn Glu
 260 265 270
 Tyr Gly Gly Asn Ile Gly Glu Thr Tyr Gly Val Pro Val Glu Glu Ile
 275 280 285

Val Glu Gly Ile Lys His Gly Val Arg Lys Val Asn Ile Asp Thr Asp
 290 295 300

Leu Arg Leu Ala Ser Thr Gly Ala Val Arg Arg Tyr Leu Ala Glu Asn
 305 310 315 320

Pro Ser Asp Phe Asp Pro Arg Lys Tyr Leu Ser Lys Thr Ile Glu Ala
 325 330 335

Met Lys Gln Ile Cys Leu Asp Arg Tyr Leu Ala Phe Gly Cys Glu Gly
 340 345 350

Gln Ala Gly Lys Ile Lys Pro Val Ser Leu Glu Lys Met Ala Asn Arg
 355 360 365

Tyr Ala Lys Gly Glu Leu Asn Gln Ile Val Lys
 370 375

<210> 127

<211> 444

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(444)

<400> 127

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 1 5 10 15

cgc cca aac aca aaa aaa att ggt aat ttt tcc tat tca agg cta caa 96
 Arg Pro Asn Thr Lys Lys Ile Gly Asn Phe Ser Tyr Ser Arg Leu Gln
 20 25 30

ttc gac acg cac agg gca tct gcc cga ttt cac aat cat aat agc gga 144
 Phe Asp Thr His Arg Ala Ser Ala Arg Phe His Asn His Asn Ser Gly
 35 40 45

gtt aaa aat atg aca gct aaa gga caa atg ttg caa gat ccc ttc ctg 192
 Val Lys Asn Met Thr Ala Lys Gly Gln Met Leu Gln Asp Pro Phe Leu
 50 55 60

aac gcc ctg cgt aaa gag cat gtt ccg gtt tcg att tac tta gtt aac 240
 Asn Ala Leu Arg Lys Glu His Val Pro Val Ser Ile Tyr Leu Val Asn

65	70	75	80	
ggg atc aaa ttg caa ggt cag gtt gag tct ttc gat caa tac gtt gtt				288
Gly Ile Lys Leu Gln Gly Gln Val Glu Ser Phe Asp Gln Tyr Val Val				
	85	90	95	
ctc ctg aga aac act tcc gtc acc caa atg gtt tac aaa cac gcc att				336
Leu Leu Arg Asn Thr Ser Val Thr Gln Met Val Tyr Lys His Ala Ile				
	100	105	110	
tcc acc atc gta ccg gca cgc tcc gtc aac cta caa cat gaa aac aga				384
Ser Thr Ile Val Pro Ala Arg Ser Val Asn Leu Gln His Glu Asn Arg				
	115	120	125	
ccc caa gcc gca ccg act tcg acc ctc gtc caa gtg gaa acc gtc cag				432
Pro Gln Ala Ala Pro Thr Ser Thr Leu Val Gln Val Glu Thr Val Gln				
	130	135	140	
cag cct gcc gaa				444
Gln Pro Ala Glu				
145				
<210> 128				
<211> 148				
<212> PRT				
<213> Neisseria meningitidis				
<400> 128				
Val Asp Ala Ala Ala Thr Leu Leu Thr Tyr Lys Cys Pro Ala Glu Asn				
1	5	10	15	
Arg Pro Asn Thr Lys Lys Ile Gly Asn Phe Ser Tyr Ser Arg Leu Gln				
	20	25	30	
Phe Asp Thr His Arg Ala Ser Ala Arg Phe His Asn His Asn Ser Gly				
	35	40	45	
Val Lys Asn Met Thr Ala Lys Gly Gln Met Leu Gln Asp Pro Phe Leu				
	50	55	60	
Asn Ala Leu Arg Lys Glu His Val Pro Val Ser Ile Tyr Leu Val Asn				
	65	70	75	80
Gly Ile Lys Leu Gln Gly Gln Val Glu Ser Phe Asp Gln Tyr Val Val				
	85	90	95	
Leu Leu Arg Asn Thr Ser Val Thr Gln Met Val Tyr Lys His Ala Ile				

100	105	110
Ser Thr Ile Val Pro Ala Arg	Ser Val Asn Leu Gln His Glu Asn Arg	
115	120	125
Pro Gln Ala Ala Pro Thr Ser Thr Leu Val Gln Val Glu Thr Val Gln		
130	135	140
Gln Pro Ala Glu		
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<210> 129
 <211> 1308
 <212> DNA
 <213> Neisseria meningitidis

<220>
 <221> CDS
 <222> (1)..(1308)

<400> 129

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Met Tyr Ala Lys Lys Gly Gly Leu Gly Leu Val Lys Ser Arg Arg Phe	
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gca cct ctt ttc gct acg cag ttt ctc ggc gcg ttc aac gac aat gtg	96
Ala Pro Leu Phe Ala Thr Gln Phe Leu Gly Ala Phe Asn Asp Asn Val	
20 25 30	
ttc aaa acc gcg ctg ttt gtg atg att ggg ttt tac ggt ttg ggg caa	144
Phe Lys Thr Ala Leu Phe Val Met Ile Gly Phe Tyr Gly Leu Gly Gln	
35 40 45	
aac ggc ttc ctg cct gcc gga cag atg ttg aac ttg ggc gcg ttg ctg	192
Asn Gly Phe Leu Pro Ala Gly Gln Met Leu Asn Leu Gly Ala Leu Leu	
50 55 60	
ttt att ttg ccg tat ttc ctg ttt tcc tcg ctg tcg ggg cag ttg ggt	240
Phe Ile Leu Pro Tyr Phe Leu Phe Ser Ser Leu Ser Gly Gln Leu Gly	
65 70 75 80	
aac aaa ttc gac aag gcc gtt ttg gcg cgt tgg gcc aag gtg ctg gaa	288
Asn Lys Phe Asp Lys Ala Val Leu Ala Arg Trp Ala Lys Val Leu Glu	
85 90 95	
atg atc att atg gcg gtg gcg gca tac ggg ttt tat atc cgg tct gcc	336

Met Ile Ile Met Ala Val Ala Ala Tyr Gly Phe Tyr Ile Arg Ser Ala	
100 105 110	
ccg ctg ctt ttg gcg tgt ctg ttt tgc atg ggc gcg caa tcg acg ctg	384
Pro Leu Leu Leu Ala Cys Leu Phe Cys Met Gly Ala Gln Ser Thr Leu	
115 120 125	
ttc ggg ccg ctg aaa tac gcc atc ctg ccc gat tat ctc gac gac aaa	432
Phe Gly Pro Leu Lys Tyr Ala Ile Leu Pro Asp Tyr Leu Asp Asp Lys	
130 135 140	
gag ttg atg atg ggc aac agc ctg att gaa tcg ggt acg ttt gtc gcc	480
Glu Leu Met Met Gly Asn Ser Leu Ile Glu Ser Gly Thr Phe Val Ala	
145 150 155 160	
atc ctg ttc ggt cag ata ctg ggg act gcg gtg gca ggt gta ccg cct	528
Ile Leu Phe Gly Gln Ile Leu Gly Thr Ala Val Ala Gly Val Pro Pro	
165 170 175	
tat att gtc ggg ata ctg gtt ttg ctg gtc gcc gta gga ggc acg gtc	576
Tyr Ile Val Gly Ile Leu Val Leu Leu Val Ala Val Gly Gly Thr Val	
180 185 190	
ggc agc ctg ttt atg ccg tcc gta ccc gcc aag gct gcc gat aca caa	624
Gly Ser Leu Phe Met Pro Ser Val Pro Ala Lys Ala Ala Asp Thr Gln	
195 200 205	
att gag tgg aat att gtc cgg ggt aca aaa tcc ctg ctg cgt gaa acg	672
Ile Glu Trp Asn Ile Val Arg Gly Thr Lys Ser Leu Leu Arg Glu Thr	
210 215 220	
gtg cgg cac aag ccc gtt ttt acc gcc att atc ggt att tcg tgg ttt	720
Val Arg His Lys Pro Val Phe Thr Ala Ile Ile Gly Ile Ser Trp Phe	
225 230 235 240	
tgg ttt gtc ggc gcg gtt tat acc acg caa ctg ccg acc ttt acc caa	768
Trp Phe Val Gly Ala Val Tyr Thr Thr Gln Leu Pro Thr Phe Thr Gln	
245 250 255	
atc cat cta ggc ggc aac gac aat gtt ttc aac ctg atg ctt gcc ctg	816
Ile His Leu Gly Gly Asn Asp Asn Val Phe Asn Leu Met Leu Ala Leu	
260 265 270	
ttt tcc atc ggt att gcc gcc ggt tcg gta ctg tgt gcc aag ttc agc	864
Phe Ser Ile Gly Ile Ala Ala Gly Ser Val Leu Cys Ala Lys Phe Ser	
275 280 285	
agg gaa cgg ctg agg ttg gct tgg gta acg gtt ggt gcg ttg ggt ttg	912

Arg Glu Arg Leu Arg Leu Ala Trp Val Thr Val Gly Ala Leu Gly Leu
 290 295 300

acg gtt tgc ggc ttg gtt ttg gtg tgg ctg acg cac gga cac cgt ttt 960
 Thr Val Cys Gly Leu Val Leu Val Trp Leu Thr His Gly His Arg Phe
 305 310 315 320

gaa ggg ctg aac ggc att ttt tgg ttt tta tcg caa gga tgg gca tat 1008
 Glu Gly Leu Asn Gly Ile Phe Trp Phe Leu Ser Gln Gly Trp Ala Tyr
 325 330 335

ccc gtg atg gcg gtg atg acg ctg atc ggc ttt ttc ggc gga ttt ttc 1056
 Pro Val Met Ala Val Met Thr Leu Ile Gly Phe Phe Gly Gly Phe Phe
 340 345 350

tcc gtt ccg ctc tat acc tgg ctg caa acc gcc agt agc gag act ttc 1104
 Ser Val Pro Leu Tyr Thr Trp Leu Gln Thr Ala Ser Ser Glu Thr Phe
 355 360 365

cgc gcc cgc gcc gtt gcc gcc aac aat atc gtt aac ggt att ttt atg 1152
 Arg Ala Arg Ala Val Ala Ala Asn Asn Ile Val Asn Gly Ile Phe Met
 370 375 380

gtt tcc gct gcc gtt ttg agc gcg gtg ttg ctg ttt ttg ttt gac agc 1200
 Val Ser Ala Ala Val Leu Ser Ala Val Leu Leu Phe Leu Phe Asp Ser
 385 390 395 400

att tcc ttg ttg tat ctg att gtc gct ttg ggc aat att ccg ttg tcg 1248
 Ile Ser Leu Leu Tyr Leu Ile Val Ala Leu Gly Asn Ile Pro Leu Ser
 405 410 415

gta ttt ttg att aag cgc gaa agg cgg ttt tta ggc gcg gcg gca atc 1296
 Val Phe Leu Ile Lys Arg Glu Arg Arg Phe Leu Gly Ala Ala Ala Ile
 420 425 430

agg aaa aaa cct 1308
 Arg Lys Lys Pro
 435

<210> 130

<211> 436

<212> PRT

<213> Neisseria meningitidis

<400> 130

Met Tyr Ala Lys Lys Gly Gly Leu Gly Leu Val Lys Ser Arg Arg Phe
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Ala Pro Leu Phe Ala Thr Gln Phe Leu Gly Ala Phe Asn Asp Asn Val
 20 25 30
 Phe Lys Thr Ala Leu Phe Val Met Ile Gly Phe Tyr Gly Leu Gly Gln
 35 40 45
 Asn Gly Phe Leu Pro Ala Gly Gln Met Leu Asn Leu Gly Ala Leu Leu
 50 55 60
 Phe Ile Leu Pro Tyr Phe Leu Phe Ser Ser Leu Ser Gly Gln Leu Gly
 65 70 75 80
 Asn Lys Phe Asp Lys Ala Val Leu Ala Arg Trp Ala Lys Val Leu Glu
 85 90 95
 Met Ile Ile Met Ala Val Ala Ala Tyr Gly Phe Tyr Ile Arg Ser Ala
 100 105 110
 Pro Leu Leu Leu Ala Cys Leu Phe Cys Met Gly Ala Gln Ser Thr Leu
 115 120 125
 Phe Gly Pro Leu Lys Tyr Ala Ile Leu Pro Asp Tyr Leu Asp Asp Lys
 130 135 140
 Glu Leu Met Met Gly Asn Ser Leu Ile Glu Ser Gly Thr Phe Val Ala
 145 150 155 160
 Ile Leu Phe Gly Gln Ile Leu Gly Thr Ala Val Ala Gly Val Pro Pro
 165 170 175
 Tyr Ile Val Gly Ile Leu Val Leu Val Ala Val Gly Gly Thr Val
 180 185 190
 Gly Ser Leu Phe Met Pro Ser Val Pro Ala Lys Ala Ala Asp Thr Gln
 195 200 205
 Ile Glu Trp Asn Ile Val Arg Gly Thr Lys Ser Leu Leu Arg Glu Thr
 210 215 220
 Val Arg His Lys Pro Val Phe Thr Ala Ile Ile Gly Ile Ser Trp Phe
 225 230 235 240
 Trp Phe Val Gly Ala Val Tyr Thr Thr Gln Leu Pro Thr Phe Thr Gln
 245 250 255
 Ile His Leu Gly Gly Asn Asp Asn Val Phe Asn Leu Met Leu Ala Leu
 260 265 270

Phe Ser Ile Gly Ile Ala Ala Gly Ser Val Leu Cys Ala Lys Phe Ser
 275 280 285
 Arg Glu Arg Leu Arg Leu Ala Trp Val Thr Val Gly Ala Leu Gly Leu
 290 295 300
 Thr Val Cys Gly Leu Val Leu Val Trp Leu Thr His Gly His Arg Phe
 305 310 315 320
 Glu Gly Leu Asn Gly Ile Phe Trp Phe Leu Ser Gln Gly Trp Ala Tyr
 325 330 335
 Pro Val Met Ala Val Met Thr Leu Ile Gly Phe Phe Gly Gly Phe Phe
 340 345 350
 Ser Val Pro Leu Tyr Thr Trp Leu Gln Thr Ala Ser Ser Glu Thr Phe
 355 360 365
 Arg Ala Arg Ala Val Ala Ala Asn Asn Ile Val Asn Gly Ile Phe Met
 370 375 380
 Val Ser Ala Ala Val Leu Ser Ala Val Leu Leu Phe Leu Phe Asp Ser
 385 390 395 400
 Ile Ser Leu Leu Tyr Leu Ile Val Ala Leu Gly Asn Ile Pro Leu Ser
 405 410 415
 Val Phe Leu Ile Lys Arg Glu Arg Arg Phe Leu Gly Ala Ala Ala Ile
 420 425 430
 Arg Lys Lys Pro
 435

<210> 131

<211> 876

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(876)

<400> 131

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1	5	10	15	
ttg aaa ggc gac atc aac gtt tcg ttt gaa ttt ttt cca ccg aaa aac				96
Leu Lys Gly Asp Ile Asn Val Ser Phe Glu Phe Phe Pro Pro Lys Asn				
20	25	30		
gag caa atg gaa acg atg ctg tgg gat tcc atc cac cgt ctg caa acc				144
Glu Gln Met Glu Thr Met Leu Trp Asp Ser Ile His Arg Leu Gln Thr				
35	40	45		
ctg cat ccc aag ttc gta tcc gta acc tac ggc gca aac tcc ggc gaa				192
Leu His Pro Lys Phe Val Ser Val Thr Tyr Gly Ala Asn Ser Gly Glu				
50	55	60		
cgc gac cgc acg cac ggc atc gtc aaa cgc atc aaa cag gaa acc ggc				240
Arg Asp Arg Thr His Gly Ile Val Lys Arg Ile Lys Gln Glu Thr Gly				
65	70	75	80	
ttg gaa gca gca ccg cac ctg acc ggc atc gac gca tcc ccc gac gaa				288
Leu Glu Ala Ala Pro His Leu Thr Gly Ile Asp Ala Ser Pro Asp Glu				
85	90	95		
ttg cgc caa atc gcc aaa gac tat tgg gac agc ggc atc cgc cgc att				336
Leu Arg Gln Ile Ala Lys Asp Tyr Trp Asp Ser Gly Ile Arg Arg Ile				
100	105	110		
gtc gcc ctg cgt ggc gac gag ccg ccc ggt tat gag aaa aaa ccg ttt				384
Val Ala Leu Arg Gly Asp Glu Pro Pro Gly Tyr Glu Lys Lys Pro Phe				
115	120	125		
tac gcc gaa gac ttg gtt aag cta tta cgc tcc gtc gcc gac ttc gac				432
Tyr Ala Glu Asp Leu Val Lys Leu Leu Arg Ser Val Ala Asp Phe Asp				
130	135	140		
atc tct gtg gcg gca tat ccc gaa gtg cat ccc gaa gcc aaa tcc gca				480
Ile Ser Val Ala Ala Tyr Pro Glu Val His Pro Glu Ala Lys Ser Ala				
145	150	155	160	
caa gcc gat ctg att aat ctg aag cgc aaa atc gat gcg ggt gca aac				528
Gln Ala Asp Leu Ile Asn Leu Lys Arg Lys Ile Asp Ala Gly Ala Asn				
165	170	175		
cac gtc atc acc caa ttt ttc ttt gac gta gaa cgc tac ctg cgc ttc				576
His Val Ile Thr Gln Phe Phe Phe Asp Val Glu Arg Tyr Leu Arg Phe				
180	185	190		
cgc gac cgc tgc gtg atg ttg ggt atc gat gtg gaa atc gtc cct ggt				624
Arg Asp Arg Cys Val Met Leu Gly Ile Asp Val Glu Ile Val Pro Gly				

195	200	205	
att ttg cct gtt acc aac ttc aag cag ctc ggc aaa atg gcg caa gta			672
Ile Leu Pro Val Thr Asn Phe Lys Gln Leu Gly Lys Met Ala Gln Val			
210	215	220	
acc aac gtc aaa atc cca agc tgg ctg tcg caa atg tat gaa ggt ttg			720
Thr Asn Val Lys Ile Pro Ser Trp Leu Ser Gln Met Tyr Glu Gly Leu			
225	230	235	240
gac gac gac caa ggc acg cgc aac ctc gtc gcc gcc agt atc gcc atc			768
Asp Asp Asp Gln Gly Thr Arg Asn Leu Val Ala Ala Ser Ile Ala Ile			
245	250	255	
gat atg gtc aaa gtc ctg tcc cgc gaa ggc gtg aaa gat ttc cac ttc			816
Asp Met Val Lys Val Leu Ser Arg Glu Gly Val Lys Asp Phe His Phe			
260	265	270	
tac acg ctc aac cgc agc gag ctg act tac gcc atc tgc cat att tta			864
Tyr Thr Leu Asn Arg Ser Glu Leu Thr Tyr Ala Ile Cys His Ile Leu			
275	280	285	
ggc gtg cgc cct			876
Gly Val Arg Pro			
290			
<210> 132			
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<400> 132			
Met Asn Tyr Ala Lys Glu Ile Asn Ala Leu Asn Asn Ser Leu Ser Asp			
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Leu Lys Gly Asp Ile Asn Val Ser Phe Glu Phe Phe Pro Pro Lys Asn			
20	25	30	
Glu Gln Met Glu Thr Met Leu Trp Asp Ser Ile His Arg Leu Gln Thr			
35	40	45	
Leu His Pro Lys Phe Val Ser Val Thr Tyr Gly Ala Asn Ser Gly Glu			
50	55	60	
Arg Asp Arg Thr His Gly Ile Val Lys Arg Ile Lys Gln Glu Thr Gly			
65	70	75	80

Leu Glu Ala Ala Pro His Leu Thr Gly Ile Asp Ala Ser Pro Asp Glu
 85 90 95

Leu Arg Gln Ile Ala Lys Asp Tyr Trp Asp Ser Gly Ile Arg Arg Ile
 100 105 110

Val Ala Leu Arg Gly Asp Glu Pro Pro Gly Tyr Glu Lys Lys Pro Phe
 115 120 125

Tyr Ala Glu Asp Leu Val Lys Leu Leu Arg Ser Val Ala Asp Phe Asp
 130 135 140

Ile Ser Val Ala Ala Tyr Pro Glu Val His Pro Glu Ala Lys Ser Ala
 145 150 155 160

Gln Ala Asp Leu Ile Asn Leu Lys Arg Lys Ile Asp Ala Gly Ala Asn
 165 170 175

His Val Ile Thr Gln Phe Phe Phe Asp Val Glu Arg Tyr Leu Arg Phe
 180 185 190

Arg Asp Arg Cys Val Met Leu Gly Ile Asp Val Glu Ile Val Pro Gly
 195 200 205

Ile Leu Pro Val Thr Asn Phe Lys Gln Leu Gly Lys Met Ala Gln Val
 210 215 220

Thr Asn Val Lys Ile Pro Ser Trp Leu Ser Gln Met Tyr Glu Gly Leu
 225 230 235 240

Asp Asp Asp Gln Gly Thr Arg Asn Leu Val Ala Ala Ser Ile Ala Ile
 245 250 255

Asp Met Val Lys Val Leu Ser Arg Glu Gly Val Lys Asp Phe His Phe
 260 265 270

Tyr Thr Leu Asn Arg Ser Glu Leu Thr Tyr Ala Ile Cys His Ile Leu
 275 280 285

Gly Val Arg Pro
 290

<210> 133

<211> 1371

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1371)

<400> 133

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  1             5             10             15

gca ctg acc gcg caa ggt tac gaa aac ccg aca ccc atc caa gcc gcc 96
Ala Leu Thr Ala Gln Gly Tyr Glu Asn Pro Thr Pro Ile Gln Ala Ala
      20             25             30

gcc att ccc aaa gca ctc gcc ggt cat gat ttg cta gcc gcc gcg caa 144
Ala Ile Pro Lys Ala Leu Ala Gly His Asp Leu Leu Ala Ala Ala Gln
      35             40             45

acc ggc aca ggc aaa acc gcc gcc ttt atg ctg ccc agt ctg gaa cgc 192
Thr Gly Thr Gly Lys Thr Ala Ala Phe Met Leu Pro Ser Leu Glu Arg
      50             55             60

ctc aaa cgt tac gcc acc gcc agc acc tcg ccc gcg atg cac ccc gtg 240
Leu Lys Arg Tyr Ala Thr Ala Ser Thr Ser Pro Ala Met His Pro Val
      65             70             75             80

cgt atg ctc gtc ctc acc ccc acg cgc gaa ctt gcc gac caa atc gac 288
Arg Met Leu Val Leu Thr Pro Thr Arg Glu Leu Ala Asp Gln Ile Asp
      85             90             95

caa aac gtg cag ggc tac atc aaa aac ctg ccg ctg cgc cac acc gtc 336
Gln Asn Val Gln Gly Tyr Ile Lys Asn Leu Pro Leu Arg His Thr Val
      100             105             110

ttg ttc ggc ggt atg aat atg gac aaa cag acc gcc gac ctg cgt gcc 384
Leu Phe Gly Gly Met Asn Met Asp Lys Gln Thr Ala Asp Leu Arg Ala
      115             120             125

ggc tgc gaa atc gtc gtc gcc acc gtc gga cgg ctg ctc gac cac gtg 432
Gly Cys Glu Ile Val Val Ala Thr Val Gly Arg Leu Leu Asp His Val
      130             135             140

aaa cag aaa aac atc cat ttg aac aaa gtc gaa atc gtc gtt ttg gac 480
Lys Gln Lys Asn Ile His Leu Asn Lys Val Glu Ile Val Val Leu Asp
      145             150             155             160

gaa gcc gac cgt atg ctg gat atg ggt ttt atc gac gac atc cgc aaa 528
Glu Ala Asp Arg Met Leu Asp Met Gly Phe Ile Asp Asp Ile Arg Lys

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	165	170	175	
atc atg cag atg ctg ccc cgc caa cgc caa acc ctg ctc ttt tcc gcc				576
Ile Met Gln Met Leu Pro Arg Gln Arg Gln Thr Leu Leu Phe Ser Ala				
	180	185	190	
acc ttc tcc gcc ccg ata cgc aaa ctg gcg caa gac ttc atg aac gcg				624
Thr Phe Ser Ala Pro Ile Arg Lys Leu Ala Gln Asp Phe Met Asn Ala				
	195	200	205	
ccc gaa acc gtc gaa gtc gcc gcg caa gac acc acc aac gcc aac gtc				672
Pro Glu Thr Val Glu Val Ala Ala Gln Asp Thr Thr Asn Ala Asn Val				
	210	215	220	
gag cag cac atc atc gcc gtc gat acc att cag aag cgc aac ctg ctc				720
Glu Gln His Ile Ile Ala Val Asp Thr Ile Gln Lys Arg Asn Leu Leu				
225	230	235	240	
gaa cgg ctg att gtc gat ttg cat atg aac cag gtc atc gtg ttc tgc				768
Glu Arg Leu Ile Val Asp Leu His Met Asn Gln Val Ile Val Phe Cys				
	245	250	255	
aaa acc aaa caa agc gtc gac cgc gta acg cgc gaa ctg gtg cgc cgc				816
Lys Thr Lys Gln Ser Val Asp Arg Val Thr Arg Glu Leu Val Arg Arg				
	260	265	270	
aac ctg tcc gca cag gcg ata cac ggc gac cgt tcc caa caa agc cgg				864
Asn Leu Ser Ala Gln Ala Ile His Gly Asp Arg Ser Gln Gln Ser Arg				
	275	280	285	
ctc gaa aca ctc aac gcc ttc aaa gac ggc aac ctg cgc gtc ctc gtc				912
Leu Glu Thr Leu Asn Ala Phe Lys Asp Gly Asn Leu Arg Val Leu Val				
	290	295	300	
gcc acc gat atc gcc gcg cgg ggg ctg gat att gcc gaa ctg ccc ttc				960
Ala Thr Asp Ile Ala Ala Arg Gly Leu Asp Ile Ala Glu Leu Pro Phe				
305	310	315	320	
gtc atc aat tac gaa atg ccc gcc cag ccc gaa gac tac gtc cac cgc				1008
Val Ile Asn Tyr Glu Met Pro Ala Gln Pro Glu Asp Tyr Val His Arg				
	325	330	335	
atc ggg cgc acg ggg cgc gcg ggc gcg gac ggc gtg gcg att tcc ctg				1056
Ile Gly Arg Thr Gly Arg Ala Gly Ala Asp Gly Val Ala Ile Ser Leu				
	340	345	350	
atg gac gaa tcc gaa cag aaa atg ttt gaa tcc att aaa gag ctg acc				1104
Met Asp Glu Ser Glu Gln Lys Met Phe Glu Ser Ile Lys Glu Leu Thr				

355	360	365	
ggc aac aag ctg ctc atc gag cgc atc gag ggc ttc gag ccg caa tgg			1152
Gly Asn Lys Leu Leu Ile Glu Arg Ile Glu Gly Phe Glu Pro Gln Trp			
370	375	380	
tgg gaa cag ggc ggc gca aaa ccg gaa aaa ccc gaa atg cgc gaa cca			1200
Trp Glu Gln Gly Gly Ala Lys Pro Glu Lys Pro Glu Met Arg Glu Pro			
385	390	395	400
aga caa cgc aac cgc tac gaa tcc gcc aaa gcg caa cgc gaa aaa aac			1248
Arg Gln Arg Asn Arg Tyr Glu Ser Ala Lys Ala Gln Arg Glu Lys Asn			
405	410	415	
acc cgg ccg gaa aat gcg gca aac gat gcg ggc gcg gct tgc gga aaa			1296
Thr Arg Pro Glu Asn Ala Ala Asn Asp Ala Gly Ala Ala Cys Gly Lys			
420	425	430	
att gcc gga cgc agc cgc cga agc cgc cgg gaa cac cgg acg tgc gcc			1344
Ile Ala Gly Arg Ser Arg Arg Ser Arg Arg Glu His Arg Thr Cys Ala			
435	440	445	
ctg ctc caa ccg cgt tac ggc gta aaa			1371
Leu Leu Gln Pro Arg Tyr Gly Val Lys			
450	455		
<210> 134			
<211> 457			
<212> PRT			
<213> Neisseria meningitidis			
<400> 134			
Met Ser Asn Pro Phe Ser Ser Leu Gly Leu Gly Thr Glu Leu Val Ser			
1	5	10	15
Ala Leu Thr Ala Gln Gly Tyr Glu Asn Pro Thr Pro Ile Gln Ala Ala			
20	25	30	
Ala Ile Pro Lys Ala Leu Ala Gly His Asp Leu Leu Ala Ala Ala Gln			
35	40	45	
Thr Gly Thr Gly Lys Thr Ala Ala Phe Met Leu Pro Ser Leu Glu Arg			
50	55	60	
Leu Lys Arg Tyr Ala Thr Ala Ser Thr Ser Pro Ala Met His Pro Val			
65	70	75	80

Arg Met Leu Val Leu Thr Pro Thr Arg Glu Leu Ala Asp Gln Ile Asp
 85 90 95

Gln Asn Val Gln Gly Tyr Ile Lys Asn Leu Pro Leu Arg His Thr Val
 100 105 110

Leu Phe Gly Gly Met Asn Met Asp Lys Gln Thr Ala Asp Leu Arg Ala
 115 120 125

Gly Cys Glu Ile Val Val Ala Thr Val Gly Arg Leu Leu Asp His Val
 130 135 140

Lys Gln Lys Asn Ile His Leu Asn Lys Val Glu Ile Val Val Leu Asp
 145 150 155 160

Glu Ala Asp Arg Met Leu Asp Met Gly Phe Ile Asp Asp Ile Arg Lys
 165 170 175

Ile Met Gln Met Leu Pro Arg Gln Arg Gln Thr Leu Leu Phe Ser Ala
 180 185 190

Thr Phe Ser Ala Pro Ile Arg Lys Leu Ala Gln Asp Phe Met Asn Ala
 195 200 205

Pro Glu Thr Val Glu Val Ala Ala Gln Asp Thr Thr Asn Ala Asn Val
 210 215 220

Glu Gln His Ile Ile Ala Val Asp Thr Ile Gln Lys Arg Asn Leu Leu
 225 230 235 240

Glu Arg Leu Ile Val Asp Leu His Met Asn Gln Val Ile Val Phe Cys
 245 250 255

Lys Thr Lys Gln Ser Val Asp Arg Val Thr Arg Glu Leu Val Arg Arg
 260 265 270

Asn Leu Ser Ala Gln Ala Ile His Gly Asp Arg Ser Gln Gln Ser Arg
 275 280 285

Leu Glu Thr Leu Asn Ala Phe Lys Asp Gly Asn Leu Arg Val Leu Val
 290 295 300

Ala Thr Asp Ile Ala Ala Arg Gly Leu Asp Ile Ala Glu Leu Pro Phe
 305 310 315 320

Val Ile Asn Tyr Glu Met Pro Ala Gln Pro Glu Asp Tyr Val His Arg
 325 330 335

Ile Gly Arg Thr Gly Arg Ala Gly Ala Asp Gly Val Ala Ile Ser Leu
 340 345 350

Met Asp Glu Ser Glu Gln Lys Met Phe Glu Ser Ile Lys Glu Leu Thr
 355 360 365

Gly Asn Lys Leu Leu Ile Glu Arg Ile Glu Gly Phe Glu Pro Gln Trp
 370 375 380

Trp Glu Gln Gly Gly Ala Lys Pro Glu Lys Pro Glu Met Arg Glu Pro
 385 390 395 400

Arg Gln Arg Asn Arg Tyr Glu Ser Ala Lys Ala Gln Arg Glu Lys Asn
 405 410 415

Thr Arg Pro Glu Asn Ala Ala Asn Asp Ala Gly Ala Ala Cys Gly Lys
 420 425 430

Ile Ala Gly Arg Ser Arg Arg Ser Arg Arg Glu His Arg Thr Cys Ala
 435 440 445

Leu Leu Gln Pro Arg Tyr Gly Val Lys
 450 455

<210> 135

<211> 1374

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1374)

<400> 135

atg aaa cac atc cac att atc ggt atc ggc ggc acg ttt atg ggt ggg 48
 Met Lys His Ile His Ile Ile Gly Ile Gly Gly Thr Phe Met Gly Gly
 1 5 10 15

att gcc gcc att gcc aaa gaa gca ggg ttt gaa gtc agc ggt tgc gat 96
 Ile Ala Ala Ile Ala Lys Glu Ala Gly Phe Glu Val Ser Gly Cys Asp
 20 25 30

gcg aag atg tat ccg ccg atg agc acc cag ctc gaa gcc ttg ggc ata 144
 Ala Lys Met Tyr Pro Pro Met Ser Thr Gln Leu Glu Ala Leu Gly Ile
 35 40 45

ggc gtg tat gaa ggc ttc gac acc gcg cag ttg gac gaa ttt aaa gcc 192
 Gly Val Tyr Glu Gly Phe Asp Thr Ala Gln Leu Asp Glu Phe Lys Ala
 50 55 60

gac gtt tac gtt atc ggc aat gtc gcc aag cgc ggg atg gat gtg gtt 240
 Asp Val Tyr Val Ile Gly Asn Val Ala Lys Arg Gly Met Asp Val Val
 65 70 75 80

gaa gcg att ttg aac cgt ggg ctg cct tat att tcc ggc ccg caa tgg 288
 Glu Ala Ile Leu Asn Arg Gly Leu Pro Tyr Ile Ser Gly Pro Gln Trp
 85 90 95

ctg gct gaa aac gtg ctg cac cat cat tgg gta ctc ggc gtg gcg ggg 336
 Leu Ala Glu Asn Val Leu His His His Trp Val Leu Gly Val Ala Gly
 100 105 110

acg cac ggc aaa acg acc acc gcg tct atg ctc gcg tgg gtt ttg gaa 384
 Thr His Gly Lys Thr Thr Thr Ala Ser Met Leu Ala Trp Val Leu Glu
 115 120 125

tat gcc gga ctc gca ccg ggc ttc ctt atc ggc ggc gta ccg gaa aac 432
 Tyr Ala Gly Leu Ala Pro Gly Phe Leu Ile Gly Gly Val Pro Glu Asn
 130 135 140

ttc agc gtt tcc gcc cgc ctg ccg caa acg ccg cgc caa gac ccg aac 480
 Phe Ser Val Ser Ala Arg Leu Pro Gln Thr Pro Arg Gln Asp Pro Asn
 145 150 155 160

agc caa tcg ccg ttt ttc gtc att gaa gcc gac gaa tac gac acc gcg 528
 Ser Gln Ser Pro Phe Phe Val Ile Glu Ala Asp Glu Tyr Asp Thr Ala
 165 170 175

ttt ttc gac aaa cgc tcc aaa ttc gtg cat tac cgt ccg cgt acc gcc 576
 Phe Phe Asp Lys Arg Ser Lys Phe Val His Tyr Arg Pro Arg Thr Ala
 180 185 190

gtg ttg aac aat ctg gaa ttc gac cac gcc gac atc ttc gcc gat ttg 624
 Val Leu Asn Asn Leu Glu Phe Asp His Ala Asp Ile Phe Ala Asp Leu
 195 200 205

ggc gcg ata cag acc cag ttc cac cac ctc gtg cgt acc gtg ccg tct 672
 Gly Ala Ile Gln Thr Gln Phe His His Leu Val Arg Thr Val Pro Ser
 210 215 220

gaa ggc ctc atc gtc tgc aac gga cgg cag caa agc ctg caa gac act 720
 Glu Gly Leu Ile Val Cys Asn Gly Arg Gln Gln Ser Leu Gln Asp Thr
 225 230 235 240

ttg gac aaa ggc tgc tgg acg ccg gtg gaa aaa ttc ggc acg gaa cac	768
Leu Asp Lys Gly Cys Trp Thr Pro Val Glu Lys Phe Gly Thr Glu His	
245 250 255	
ggc tgg cag gcc ggc gaa gcc aat gcc gat ggc tcg ttc gac gtg ttg	816
Gly Trp Gln Ala Gly Glu Ala Asn Ala Asp Gly Ser Phe Asp Val Leu	
260 265 270	
ctt gac ggc aaa aaa gcc gga cac gtc gct tgg agt ttg atg ggc gga	864
Leu Asp Gly Lys Lys Ala Gly His Val Ala Trp Ser Leu Met Gly Gly	
275 280 285	
cac aac cgc atg aac gcg ctc gcc gtc atc gcc gcc gcg cgt cat gcc	912
His Asn Arg Met Asn Ala Leu Ala Val Ile Ala Ala Ala Arg His Ala	
290 295 300	
gga gtc gac att cag acg gcc tgc gaa gcc ttg agc acg ttt aaa aac	960
Gly Val Asp Ile Gln Thr Ala Cys Glu Ala Leu Ser Thr Phe Lys Asn	
305 310 315 320	
gtc aaa cgc cgc atg gaa atc aaa ggc acg gca aac ggt atc acc gtt	1008
Val Lys Arg Arg Met Glu Ile Lys Gly Thr Ala Asn Gly Ile Thr Val	
325 330 335	
tac gac gac ttc gcc cac cat ccg acc gct atc gaa acc acg att caa	1056
Tyr Asp Asp Phe Ala His His Pro Thr Ala Ile Glu Thr Thr Ile Gln	
340 345 350	
ggt ttg cgc cag cgc gtc ggc ggc gcg cgc atc ctc gcc gtc ctc gaa	1104
Gly Leu Arg Gln Arg Val Gly Gly Ala Arg Ile Leu Ala Val Leu Glu	
355 360 365	
ccg cgt tcc aat acg atg aag ctg ggt acg atg aaa gcc gcc ctg ccc	1152
Pro Arg Ser Asn Thr Met Lys Leu Gly Thr Met Lys Ala Ala Leu Pro	
370 375 380	
gca agc ctc aaa gaa gcc gac caa gtg ttc tgc tac gcc ggc ggc gcg	1200
Ala Ser Leu Lys Glu Ala Asp Gln Val Phe Cys Tyr Ala Gly Gly Ala	
385 390 395 400	
gac tgg gac gtt gcc gaa gcc ctc gcg cct ttg ggc ggc agg ctg cac	1248
Asp Trp Asp Val Ala Glu Ala Leu Ala Pro Leu Gly Gly Arg Leu His	
405 410 415	
gtc ggc aaa gac ttc gat gcc ttc gtt gcc gaa atc gtg aaa aac gcc	1296
Val Gly Lys Asp Phe Asp Ala Phe Val Ala Glu Ile Val Lys Asn Ala	
420 425 430	

gaa gca ggc gac cat att ttg gtg atg agc aac ggc ggt ttc ggc gga 1344
 Glu Ala Gly Asp His Ile Leu Val Met Ser Asn Gly Gly Phe Gly Gly
 435 440 445

ata cac acc aaa ctg ctg gac gct ttg aga 1374
 Ile His Thr Lys Leu Leu Asp Ala Leu Arg
 450 455

<210> 136

<211> 458

<212> PRT

<213> Neisseria meningitidis

<400> 136

Met Lys His Ile His Ile Ile Gly Ile Gly Gly Thr Phe Met Gly Gly
 1 5 10 15

Ile Ala Ala Ile Ala Lys Glu Ala Gly Phe Glu Val Ser Gly Cys Asp
 20 25 30

Ala Lys Met Tyr Pro Pro Met Ser Thr Gln Leu Glu Ala Leu Gly Ile
 35 40 45

Gly Val Tyr Glu Gly Phe Asp Thr Ala Gln Leu Asp Glu Phe Lys Ala
 50 55 60

Asp Val Tyr Val Ile Gly Asn Val Ala Lys Arg Gly Met Asp Val Val
 65 70 75 80

Glu Ala Ile Leu Asn Arg Gly Leu Pro Tyr Ile Ser Gly Pro Gln Trp
 85 90 95

Leu Ala Glu Asn Val Leu His His His Trp Val Leu Gly Val Ala Gly
 100 105 110

Thr His Gly Lys Thr Thr Thr Ala Ser Met Leu Ala Trp Val Leu Glu
 115 120 125

Tyr Ala Gly Leu Ala Pro Gly Phe Leu Ile Gly Gly Val Pro Glu Asn
 130 135 140

Phe Ser Val Ser Ala Arg Leu Pro Gln Thr Pro Arg Gln Asp Pro Asn
 145 150 155 160

Ser Gln Ser Pro Phe Phe Val Ile Glu Ala Asp Glu Tyr Asp Thr Ala
 165 170 175

Phe Phe Asp Lys Arg Ser Lys Phe Val His Tyr Arg Pro Arg Thr Ala
 180 185 190

Val Leu Asn Asn Leu Glu Phe Asp His Ala Asp Ile Phe Ala Asp Leu
 195 200 205

Gly Ala Ile Gln Thr Gln Phe His His Leu Val Arg Thr Val Pro Ser
 210 215 220

Glu Gly Leu Ile Val Cys Asn Gly Arg Gln Gln Ser Leu Gln Asp Thr
 225 230 235 240

Leu Asp Lys Gly Cys Trp Thr Pro Val Glu Lys Phe Gly Thr Glu His
 245 250 255

Gly Trp Gln Ala Gly Glu Ala Asn Ala Asp Gly Ser Phe Asp Val Leu
 260 265 270

Leu Asp Gly Lys Lys Ala Gly His Val Ala Trp Ser Leu Met Gly Gly
 275 280 285

His Asn Arg Met Asn Ala Leu Ala Val Ile Ala Ala Ala Arg His Ala
 290 295 300

Gly Val Asp Ile Gln Thr Ala Cys Glu Ala Leu Ser Thr Phe Lys Asn
 305 310 315 320

Val Lys Arg Arg Met Glu Ile Lys Gly Thr Ala Asn Gly Ile Thr Val
 325 330 335

Tyr Asp Asp Phe Ala His His Pro Thr Ala Ile Glu Thr Thr Ile Gln
 340 345 350

Gly Leu Arg Gln Arg Val Gly Gly Ala Arg Ile Leu Ala Val Leu Glu
 355 360 365

Pro Arg Ser Asn Thr Met Lys Leu Gly Thr Met Lys Ala Ala Leu Pro
 370 375 380

Ala Ser Leu Lys Glu Ala Asp Gln Val Phe Cys Tyr Ala Gly Gly Ala
 385 390 395 400

Asp Trp Asp Val Ala Glu Ala Leu Ala Pro Leu Gly Gly Arg Leu His
 405 410 415

Val Gly Lys Asp Phe Asp Ala Phe Val Ala Glu Ile Val Lys Asn Ala
 420 425 430

Glu Ala Gly Asp His Ile Leu Val Met Ser Asn Gly Gly Phe Gly Gly
 435 440 445

Ile His Thr Lys Leu Leu Asp Ala Leu Arg
 450 455

<210> 137

<211> 876

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(876)

<400> 137

ttg acc gtc cga acg aaa aag acg gcg cat tat acc cta ttc cat tcc 48
 Leu Thr Val Arg Thr Lys Lys Thr Ala His Tyr Thr Leu Phe His Ser
 1 5 10 15

gac cga aaa ccg aac atg act act ctc aaa ccc gcc ctg ccc gct tat 96
 Asp Arg Lys Pro Asn Met Thr Thr Leu Lys Pro Ala Leu Pro Ala Tyr
 20 25 30

ctg gac aac atc cgc atc atc ctc acg cgc acc agc cat ccc gcc aac 144
 Leu Asp Asn Ile Arg Ile Ile Leu Thr Arg Thr Ser His Pro Ala Asn
 35 40 45

atc ggc tct gcc gcg cgc gcg atg aaa aca atg ggt ctg cac aaa ctg 192
 Ile Gly Ser Ala Ala Arg Ala Met Lys Thr Met Gly Leu His Lys Leu
 50 55 60

acc atc gtc gcc cca aat ctg atg gca acg ccg atg acg gaa aac ccg 240
 Thr Ile Val Ala Pro Asn Leu Met Ala Thr Pro Met Thr Glu Asn Pro
 65 70 75 80

ccc gtg ttt gac ccg gag cat cct caa tcg ttt aaa tta ccg gaa gaa 288
 Pro Val Phe Asp Pro Glu His Pro Gln Ser Phe Lys Leu Pro Glu Glu
 85 90 95

agc ttc atc ctc gct tcc ggc gcg gca gac gtt ttg gaa aat gcc acc 336
 Ser Phe Ile Leu Ala Ser Gly Ala Ala Asp Val Leu Glu Asn Ala Thr
 100 105 110

att gcc gct tct ttg gac gaa gcc ctt gcc gac acc acc atc gcc tgc 384
 Ile Ala Ala Ser Leu Asp Glu Ala Leu Ala Asp Thr Thr Ile Ala Cys

115	120	125	
gcc ctg acc agc cgc cgc cgc gaa att act gcg ccg ctg caa acc ccg			432
Ala Leu Thr Ser Arg Arg Arg Glu Ile Thr Ala Pro Leu Gln Thr Pro			
130	135	140	
cgc gat ttg gta tcc gaa tta ctg cag acc gca aac cga ggc gag aaa			480
Arg Asp Leu Val Ser Glu Leu Leu Gln Thr Ala Asn Arg Gly Glu Lys			
145	150	155	160
gtg gca ctg gtt ttc ggc aac gag act ttc ggc ttg agc atc gaa gaa			528
Val Ala Leu Val Phe Gly Asn Glu Thr Phe Gly Leu Ser Ile Glu Glu			
165	170	175	
gtc caa gcc tgc aac cga ctg atg acc atc aac ggc aat ccc gac tat			576
Val Gln Ala Cys Asn Arg Leu Met Thr Ile Asn Gly Asn Pro Asp Tyr			
180	185	190	
ttc tcg ctc aac ctc gcc caa gcc gtg cag gtc gtg tgc tac gaa atc			624
Phe Ser Leu Asn Leu Ala Gln Ala Val Gln Val Val Cys Tyr Glu Ile			
195	200	205	
ttc agc caa acc ggt tcg ccc atg acc cat ctt caa caa gaa gac cac			672
Phe Ser Gln Thr Gly Ser Pro Met Thr His Leu Gln Gln Glu Asp His			
210	215	220	
gct gcg acc cac gag caa atc aaa ggc atg gtc gcc cac atg gaa agc			720
Ala Ala Thr His Glu Gln Ile Lys Gly Met Val Ala His Met Glu Ser			
225	230	235	240
gtg atg aac gac atc ggc ttt ttc aac cgc cgc aac ggc gag cgt ctg			768
Val Met Asn Asp Ile Gly Phe Phe Asn Arg Arg Asn Gly Glu Arg Leu			
245	250	255	
atg cgc cgt atg cag agc ctg ttc gga cgc gcc aac acg caa acc gaa			816
Met Arg Arg Met Gln Ser Leu Phe Gly Arg Ala Asn Thr Gln Thr Glu			
260	265	270	
gac atc gat atc ctg cgc ggt ttt ttc aat acc gtc agc cat cgt atc			864
Asp Ile Asp Ile Leu Arg Gly Phe Phe Asn Thr Val Ser His Arg Ile			
275	280	285	
cat aaa aaa gac			876
His Lys Lys Asp			
290			

<211> 292

<212> PRT

<213> Neisseria meningitidis

<400> 138

Leu Thr Val Arg Thr Lys Lys Thr Ala His Tyr Thr Leu Phe His Ser
 1 5 10 15

Asp Arg Lys Pro Asn Met Thr Thr Leu Lys Pro Ala Leu Pro Ala Tyr
 20 25 30

Leu Asp Asn Ile Arg Ile Ile Leu Thr Arg Thr Ser His Pro Ala Asn
 35 40 45

Ile Gly Ser Ala Ala Arg Ala Met Lys Thr Met Gly Leu His Lys Leu
 50 55 60

Thr Ile Val Ala Pro Asn Leu Met Ala Thr Pro Met Thr Glu Asn Pro
 65 70 75 80

Pro Val Phe Asp Pro Glu His Pro Gln Ser Phe Lys Leu Pro Glu Glu
 85 90 95

Ser Phe Ile Leu Ala Ser Gly Ala Ala Asp Val Leu Glu Asn Ala Thr
 100 105 110

Ile Ala Ala Ser Leu Asp Glu Ala Leu Ala Asp Thr Thr Ile Ala Cys
 115 120 125

Ala Leu Thr Ser Arg Arg Arg Glu Ile Thr Ala Pro Leu Gln Thr Pro
 130 135 140

Arg Asp Leu Val Ser Glu Leu Leu Gln Thr Ala Asn Arg Gly Glu Lys
 145 150 155 160

Val Ala Leu Val Phe Gly Asn Glu Thr Phe Gly Leu Ser Ile Glu Glu
 165 170 175

Val Gln Ala Cys Asn Arg Leu Met Thr Ile Asn Gly Asn Pro Asp Tyr
 180 185 190

Phe Ser Leu Asn Leu Ala Gln Ala Val Gln Val Val Cys Tyr Glu Ile
 195 200 205

Phe Ser Gln Thr Gly Ser Pro Met Thr His Leu Gln Gln Glu Asp His
 210 215 220

Ala Ala Thr His Glu Gln Ile Lys Gly Met Val Ala His Met Glu Ser

225 230 235 240
 Val Met Asn Asp Ile Gly Phe Phe Asn Arg Arg Asn Gly Glu Arg Leu
 245 250 255
 Met Arg Arg Met Gln Ser Leu Phe Gly Arg Ala Asn Thr Gln Thr Glu
 260 265 270
 Asp Ile Asp Ile Leu Arg Gly Phe Phe Asn Thr Val Ser His Arg Ile
 275 280 285
 His Lys Lys Asp
 290

<210> 139
 <211> 708
 <212> DNA
 <213> Neisseria meningitidis

<220>
 <221> CDS
 <222> (1)..(708)

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 Leu Arg Gln Leu Gly Cys Leu Ile Gly Phe Phe Ser Val Gly Ile Ile
 1 5 10 15
 atg aat ttg aaa tta gtg ttt gaa tcg ggc gat ccc gtc ctg atc ggt 96
 Met Asn Leu Lys Leu Val Phe Glu Ser Gly Asp Pro Val Leu Ile Gly
 20 25 30
 gtg ttt gtg ttg atg ctg ttg atg agt atc gta acg tgg tgt ttg gtt 144
 Val Phe Val Leu Met Leu Leu Met Ser Ile Val Thr Trp Cys Leu Val
 35 40 45
 gtc ttg cgc tgc atc aag ctg tat cgg gcg cgc aaa ggg aat gcc gcc 192
 Val Leu Arg Cys Ile Lys Leu Tyr Arg Ala Arg Lys Gly Asn Ala Ala
 50 55 60
 gtc aaa cgg cat atg cgc gat act ttg tcg ctg aac gac gcg gtc gaa 240
 Val Lys Arg His Met Arg Asp Thr Leu Ser Leu Asn Asp Ala Val Glu
 65 70 75 80
 aaa gtg cgc gcc gtc gat gcg cct ttg tcc aaa ctg gcg caa gag gca 288
 Lys Val Arg Ala Val Asp Ala Pro Leu Ser Lys Leu Ala Gln Glu Ala

85	90	95	
ttg cag tct tac cgc aac tac cgc cga aac gaa gcg tcc gaa ctg gcg			336
Leu Gln Ser Tyr Arg Asn Tyr Arg Arg Asn Glu Ala Ser Glu Leu Ala			
100	105	110	
cag gct ttg ccg ttg aac gag tat ttg gtc att caa atc cgc aac agt			384
Gln Ala Leu Pro Leu Asn Glu Tyr Leu Val Ile Gln Ile Arg Asn Ser			
115	120	125	
atg gcg cag att atg cgc cgg ttt gat tac ggg atg acc gcg ctt gcc			432
Met Ala Gln Ile Met Arg Arg Phe Asp Tyr Gly Met Thr Ala Leu Ala			
130	135	140	
tcc atc ggc gcg acc gcg ccg ttt atc ggg ttg ttc ggc acg gtt tgg			480
Ser Ile Gly Ala Thr Ala Pro Phe Ile Gly Leu Phe Gly Thr Val Trp			
145	150	155	160
ggg att tac cac gcc ctg atc aat atc ggg caa agc ggg cag atg agt			528
Gly Ile Tyr His Ala Leu Ile Asn Ile Gly Gln Ser Gly Gln Met Ser			
165	170	175	
att gcg gcg gtt gcc ggc ccg att ggc gag gca ctg gtg gcg acg gcg			576
Ile Ala Ala Val Ala Gly Pro Ile Gly Glu Ala Leu Val Ala Thr Ala			
180	185	190	
gcg ggt ttg ttc gtg gcg att ccg gcg gtg ttg gca tac aac ttc ctc			624
Ala Gly Leu Phe Val Ala Ile Pro Ala Val Leu Ala Tyr Asn Phe Leu			
195	200	205	
aat cgc ggc aca aaa ata ctg acc cag gat ttg gat gcg atg gcg cac			672
Asn Arg Gly Thr Lys Ile Leu Thr Gln Asp Leu Asp Ala Met Ala His			
210	215	220	
gat ttg cac gtc cgc ctg ctt aat caa aag gat agc			708
Asp Leu His Val Arg Leu Leu Asn Gln Lys Asp Ser			
225	230	235	
<210> 140			
<211> 236			
<212> PRT			
<213> Neisseria meningitidis			
<400> 140			
Leu Arg Gln Leu Gly Cys Leu Ile Gly Phe Phe Ser Val Gly Ile Ile			
1	5	10	15

Met Asn Leu Lys Leu Val Phe Glu Ser Gly Asp Pro Val Leu Ile Gly
 20 25 30
 Val Phe Val Leu Met Leu Leu Met Ser Ile Val Thr Trp Cys Leu Val
 35 40 45
 Val Leu Arg Cys Ile Lys Leu Tyr Arg Ala Arg Lys Gly Asn Ala Ala
 50 55 60
 Val Lys Arg His Met Arg Asp Thr Leu Ser Leu Asn Asp Ala Val Glu
 65 70 75 80
 Lys Val Arg Ala Val Asp Ala Pro Leu Ser Lys Leu Ala Gln Glu Ala
 85 90 95
 Leu Gln Ser Tyr Arg Asn Tyr Arg Arg Asn Glu Ala Ser Glu Leu Ala
 100 105 110
 Gln Ala Leu Pro Leu Asn Glu Tyr Leu Val Ile Gln Ile Arg Asn Ser
 115 120 125
 Met Ala Gln Ile Met Arg Arg Phe Asp Tyr Gly Met Thr Ala Leu Ala
 130 135 140
 Ser Ile Gly Ala Thr Ala Pro Phe Ile Gly Leu Phe Gly Thr Val Trp
 145 150 155 160
 Gly Ile Tyr His Ala Leu Ile Asn Ile Gly Gln Ser Gly Gln Met Ser
 165 170 175
 Ile Ala Ala Val Ala Gly Pro Ile Gly Glu Ala Leu Val Ala Thr Ala
 180 185 190
 Ala Gly Leu Phe Val Ala Ile Pro Ala Val Leu Ala Tyr Asn Phe Leu
 195 200 205
 Asn Arg Gly Thr Lys Ile Leu Thr Gln Asp Leu Asp Ala Met Ala His
 210 215 220
 Asp Leu His Val Arg Leu Leu Asn Gln Lys Asp Ser
 225 230 235

<210> 141

<211> 812

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(810)

<400> 141

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atg act atg cac gcc ctc ccc cgc tac gcc gtt ttt ggc aac ccc gtc 48
Met Thr Met His Ala Leu Pro Arg Tyr Ala Val Phe Gly Asn Pro Val
  1             5             10             15

gcc cac agc aaa tcg ccg caa att cat caa caa ttt gcc ctt cag gaa 96
Ala His Ser Lys Ser Pro Gln Ile His Gln Gln Phe Ala Leu Gln Glu
      20             25             30

ggc gtt gac att gaa tac gaa cgc att tgc gcc gac atc ggc ggt ttc 144
Gly Val Asp Ile Glu Tyr Glu Arg Ile Cys Ala Asp Ile Gly Gly Phe
      35             40             45

gcg cag gcg gtt tcg aca ttt ttt gaa aca ggc ggt tgc ggg gca aac 192
Ala Gln Ala Val Ser Thr Phe Phe Glu Thr Gly Gly Cys Gly Ala Asn
      50             55             60

gtt acc gta ccg ttc aag cag gaa gcg ttt cat ctg gcg gac gag cat 240
Val Thr Val Pro Phe Lys Gln Glu Ala Phe His Leu Ala Asp Glu His
      65             70             75             80

tct gaa cgc gca ttg gct gcc ggc gcg gtc aac acg ctg att ttt ctg 288
Ser Glu Arg Ala Leu Ala Ala Gly Ala Val Asn Thr Leu Ile Phe Leu
      85             90             95

aaa aac gga aaa ctg cgc ggc gac aat acc gac ggt atc ggt ttg gcc 336
Lys Asn Gly Lys Leu Arg Gly Asp Asn Thr Asp Gly Ile Gly Leu Ala
      100             105             110

aac gac atc acg cag gtc aaa aac att gcc atc gaa ggc aaa acc atc 384
Asn Asp Ile Thr Gln Val Lys Asn Ile Ala Ile Glu Gly Lys Thr Ile
      115             120             125

ttg ctt ttg ggc gcg ggc ggc gcg gtg cgc ggc gtg att cct gtt ttg 432
Leu Leu Leu Gly Ala Gly Gly Ala Val Arg Gly Val Ile Pro Val Leu
      130             135             140

aaa gaa cac cgt cct gcc cgt atc gtc att gcc aac cgt acc cgc gcc 480
Lys Glu His Arg Pro Ala Arg Ile Val Ile Ala Asn Arg Thr Arg Ala
      145             150             155             160

aaa gcc gag gaa ttg gcg cag ctt ttc ggc att gaa gcc gtc ccg atg 528
Lys Ala Glu Glu Leu Ala Gln Leu Phe Gly Ile Glu Ala Val Pro Met

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165	170	175	
gcg gat gtg aac ggc ggt ttt gat atc atc atc aac ggc acg tcg ggc	576		
Ala Asp Val Asn Gly Gly Phe Asp Ile Ile Ile Asn Gly Thr Ser Gly			
180	185	190	
ggt cta aac ggt cag att ccc gat att ccg ccc gat att ttt caa aac	624		
Gly Leu Asn Gly Gln Ile Pro Asp Ile Pro Pro Asp Ile Phe Gln Asn			
195	200	205	
tgc gcg ctt gcc tac gat atg gtg tac ggc tgc gcg gca aaa ccg ttt	672		
Cys Ala Leu Ala Tyr Asp Met Val Tyr Gly Cys Ala Ala Lys Pro Phe			
210	215	220	
tta gat ttt gca cga caa tcg ggt gcg aaa aaa act gcc gac gga ctg	720		
Leu Asp Phe Ala Arg Gln Ser Gly Ala Lys Lys Thr Ala Asp Gly Leu			
225	230	235	240
ggt atg cta gtc ggt caa gcg gcg gct tcc tac gcc ctc tgg cgc gga	768		
Gly Met Leu Val Gly Gln Ala Ala Ala Ser Tyr Ala Leu Trp Arg Gly			
245	250	255	
ttt acg ccc gat atc cgc ccc gtt atc gaa tac atg aaa gcc ct	812		
Phe Thr Pro Asp Ile Arg Pro Val Ile Glu Tyr Met Lys Ala			
260	265	270	
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<211> 270			
<212> PRT			
<213> Neisseria meningitidis			
<400> 142			
Met Thr Met His Ala Leu Pro Arg Tyr Ala Val Phe Gly Asn Pro Val			
1	5	10	15
Ala His Ser Lys Ser Pro Gln Ile His Gln Gln Phe Ala Leu Gln Glu			
20	25	30	
Gly Val Asp Ile Glu Tyr Glu Arg Ile Cys Ala Asp Ile Gly Gly Phe			
35	40	45	
Ala Gln Ala Val Ser Thr Phe Phe Glu Thr Gly Gly Cys Gly Ala Asn			
50	55	60	
Val Thr Val Pro Phe Lys Gln Glu Ala Phe His Leu Ala Asp Glu His			
65	70	75	80

Ser Glu Arg Ala Leu Ala Ala Gly Ala Val Asn Thr Leu Ile Phe Leu
 85 90 95

Lys Asn Gly Lys Leu Arg Gly Asp Asn Thr Asp Gly Ile Gly Leu Ala
 100 105 110

Asn Asp Ile Thr Gln Val Lys Asn Ile Ala Ile Glu Gly Lys Thr Ile
 115 120 125

Leu Leu Leu Gly Ala Gly Gly Ala Val Arg Gly Val Ile Pro Val Leu
 130 135 140

Lys Glu His Arg Pro Ala Arg Ile Val Ile Ala Asn Arg Thr Arg Ala
 145 150 155 160

Lys Ala Glu Glu Leu Ala Gln Leu Phe Gly Ile Glu Ala Val Pro Met
 165 170 175

Ala Asp Val Asn Gly Gly Phe Asp Ile Ile Ile Asn Gly Thr Ser Gly
 180 185 190

Gly Leu Asn Gly Gln Ile Pro Asp Ile Pro Pro Asp Ile Phe Gln Asn
 195 200 205

Cys Ala Leu Ala Tyr Asp Met Val Tyr Gly Cys Ala Ala Lys Pro Phe
 210 215 220

Leu Asp Phe Ala Arg Gln Ser Gly Ala Lys Lys Thr Ala Asp Gly Leu
 225 230 235 240

Gly Met Leu Val Gly Gln Ala Ala Ala Ser Tyr Ala Leu Trp Arg Gly
 245 250 255

Phe Thr Pro Asp Ile Arg Pro Val Ile Glu Tyr Met Lys Ala
 260 265 270

<210> 143

<211> 1515

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1515)

<400> 143

atg ttg tat ttc aga tac ggt ttt ttg gtt gtt tgg tgt gcg gca ggt	48
Met Leu Tyr Phe Arg Tyr Gly Phe Leu Val Val Trp Cys Ala Ala Gly	
1 5 10 15	
gtt tct gcc gcc tat ggg gcg gat gcg ccc gcg att ttg gat gac aag	96
Val Ser Ala Ala Tyr Gly Ala Asp Ala Pro Ala Ile Leu Asp Asp Lys	
20 25 30	
gca ttg ttg cag gtg cag cgg tcg gtg tcg gat aag tgg gcg gaa tcg	144
Ala Leu Leu Gln Val Gln Arg Ser Val Ser Asp Lys Trp Ala Glu Ser	
35 40 45	
gat tgg aaa gtt gac aat gat gcc ccg cgc gtg gtt gac ggg gat ttt	192
Asp Trp Lys Val Asp Asn Asp Ala Pro Arg Val Val Asp Gly Asp Phe	
50 55 60	
ttg ttg gcg cat ccg aaa atg ttg gaa cat agt ttg cgc gac gtg ctc	240
Leu Leu Ala His Pro Lys Met Leu Glu His Ser Leu Arg Asp Val Leu	
65 70 75 80	
aac ggc aat cag gcg gat ttg atc gct tcg ttg gcg gat ttg tat gcc	288
Asn Gly Asn Gln Ala Asp Leu Ile Ala Ser Leu Ala Asp Leu Tyr Ala	
85 90 95	
aag ctg ccg gat tat gac gcg gtt ttg tac ggc agg gcg cgg gct ttg	336
Lys Leu Pro Asp Tyr Asp Ala Val Leu Tyr Gly Arg Ala Arg Ala Leu	
100 105 110	
ctg gcg aaa ttg gcg gga agg ccg gcg gag gcg gtg gcg cgg tat cgg	384
Leu Ala Lys Leu Ala Gly Arg Pro Ala Glu Ala Val Ala Arg Tyr Arg	
115 120 125	
gaa ctg cac ggg gaa aat gcg gca gac gag cgg att ttg ctg gat ttg	432
Glu Leu His Gly Glu Asn Ala Ala Asp Glu Arg Ile Leu Leu Asp Leu	
130 135 140	
gcg gcg gcg gag ttt gac gat ttc ccg ctg aag tcg gca gaa agg cat	480
Ala Ala Ala Glu Phe Asp Asp Phe Arg Leu Lys Ser Ala Glu Arg His	
145 150 155 160	
ttt gcc gag gcg gaa aaa ttg gat ttg ccg gcg ccg gtt ttg gaa aat	528
Phe Ala Glu Ala Glu Lys Leu Asp Leu Pro Ala Pro Val Leu Glu Asn	
165 170 175	
gtg ggg cgt ttt ccg aaa aaa gcg gag ggg ctg acg ggc tgg cgt ttt	576
Val Gly Arg Phe Arg Lys Lys Ala Glu Gly Leu Thr Gly Trp Arg Phe	
180 185 190	

tcg ggc ggc atc agt ccg gcg gtc aat aga aat gcc aat aat gcc gcg	624
Ser Gly Gly Ile Ser Pro Ala Val Asn Arg Asn Ala Asn Asn Ala Ala	
195 200 205	
ccg cag tat tgc cgg caa aac gga ggc cgg cag ata tgc agt gtc agc	672
Pro Gln Tyr Cys Arg Gln Asn Gly Gly Arg Gln Ile Cys Ser Val Ser	
210 215 220	
cgg gcg gag cgg gcg gca ggc ttg aat tat gaa atc gag gcg gaa aaa	720
Arg Ala Glu Arg Ala Ala Gly Leu Asn Tyr Glu Ile Glu Ala Glu Lys	
225 230 235 240	
ctg acg gcg ttg gca gat aat cat tat ttg ttg ttc cgt tcc aat atc	768
Leu Thr Ala Leu Ala Asp Asn His Tyr Leu Leu Phe Arg Ser Asn Ile	
245 250 255	
ggc ggc acg agc tat tat ttc agt aaa aaa tca gct tat gac gac ggg	816
Gly Gly Thr Ser Tyr Tyr Phe Ser Lys Lys Ser Ala Tyr Asp Asp Gly	
260 265 270	
ttc ggc aga gcg tat ttg ggt tgg cag tat aaa aat gca cgg cag acg	864
Phe Gly Arg Ala Tyr Leu Gly Trp Gln Tyr Lys Asn Ala Arg Gln Thr	
275 280 285	
gcg ggg att ttg ccg ttt tat cag gtg cag ttg tcg ggc agc gac ggc	912
Ala Gly Ile Leu Pro Phe Tyr Gln Val Gln Leu Ser Gly Ser Asp Gly	
290 295 300	
ttt gat gcg aaa aca aaa cgg gta aac aac cgc cgc ctg ccg ccg tat	960
Phe Asp Ala Lys Thr Lys Arg Val Asn Asn Arg Arg Leu Pro Pro Tyr	
305 310 315 320	
atg ctg gcg cac gga gtc ggc gtg cag ttg tcc cat act tac cgc cca	1008
Met Leu Ala His Gly Val Gly Val Gln Leu Ser His Thr Tyr Arg Pro	
325 330 335	
aac ccg gga tgg caa ttt tcg gtc gcg ctg gaa cat tac cgc caa cgc	1056
Asn Pro Gly Trp Gln Phe Ser Val Ala Leu Glu His Tyr Arg Gln Arg	
340 345 350	
tac cgc gaa cag gat agg gcg gaa tac aat aac ggt cgg cag gac ggg	1104
Tyr Arg Glu Gln Asp Arg Ala Glu Tyr Asn Asn Gly Arg Gln Asp Gly	
355 360 365	
ttt tat gtt tcg tcg gca aaa cgt ttg ggc gaa tcg gca act gtg ttc	1152
Phe Tyr Val Ser Ser Ala Lys Arg Leu Gly Glu Ser Ala Thr Val Phe	
370 375 380	

ggc ggc tgg cag ttt gtg cgg ttt gtg ccg aaa cgc gaa acg gtg ggc 1200
 Gly Gly Trp Gln Phe Val Arg Phe Val Pro Lys Arg Glu Thr Val Gly
 385 390 395 400

ggc gcg gtc aat aat gcc gcc tac cgg cgc aac ggt gtt tat gcc ggc 1248
 Gly Ala Val Asn Asn Ala Ala Tyr Arg Arg Asn Gly Val Tyr Ala Gly
 405 410 415

tgg gcg cag gag tgg cgg cag ttg ggc ggt ttg aac agt cgg gtt tcc 1296
 Trp Ala Gln Glu Trp Arg Gln Leu Gly Gly Leu Asn Ser Arg Val Ser
 420 425 430

gcg tct tat gcc cgc cgc aac tat aag ggc gtt gcg gct ttc tcg aca 1344
 Ala Ser Tyr Ala Arg Arg Asn Tyr Lys Gly Val Ala Ala Phe Ser Thr
 435 440 445

gag gcg caa cgc aac cgc gaa tgg aat gtc tcg ctg gct ttg agc cac 1392
 Glu Ala Gln Arg Asn Arg Glu Trp Asn Val Ser Leu Ala Leu Ser His
 450 455 460

gac aag ttg tcg tac aaa ggt atc gtg ccc gcg ttg aat tat cgt ttc 1440
 Asp Lys Leu Ser Tyr Lys Gly Ile Val Pro Ala Leu Asn Tyr Arg Phe
 465 470 475 480

ggc agg acg gaa agt aat gtg ccg tat gcg aaa cgc cgc aac agc gag 1488
 Gly Arg Thr Glu Ser Asn Val Pro Tyr Ala Lys Arg Arg Asn Ser Glu
 485 490 495

gtg ttt gtg tcg gcg gat tgg cgg ttt 1515
 Val Phe Val Ser Ala Asp Trp Arg Phe
 500 505

<210> 144

<211> 505

<212> PRT

<213> Neisseria meningitidis

<400> 144

Met Leu Tyr Phe Arg Tyr Gly Phe Leu Val Val Trp Cys Ala Ala Gly
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Val Ser Ala Ala Tyr Gly Ala Asp Ala Pro Ala Ile Leu Asp Asp Lys
 20 25 30

Ala Leu Leu Gln Val Gln Arg Ser Val Ser Asp Lys Trp Ala Glu Ser
 35 40 45

Asp Trp Lys Val Asp Asn Asp Ala Pro Arg Val Val Asp Gly Asp Phe
 50 55 60

Leu Leu Ala His Pro Lys Met Leu Glu His Ser Leu Arg Asp Val Leu
 65 70 75 80

Asn Gly Asn Gln Ala Asp Leu Ile Ala Ser Leu Ala Asp Leu Tyr Ala
 85 90 95

Lys Leu Pro Asp Tyr Asp Ala Val Leu Tyr Gly Arg Ala Arg Ala Leu
 100 105 110

Leu Ala Lys Leu Ala Gly Arg Pro Ala Glu Ala Val Ala Arg Tyr Arg
 115 120 125

Glu Leu His Gly Glu Asn Ala Ala Asp Glu Arg Ile Leu Leu Asp Leu
 130 135 140

Ala Ala Ala Glu Phe Asp Asp Phe Arg Leu Lys Ser Ala Glu Arg His
 145 150 155 160

Phe Ala Glu Ala Glu Lys Leu Asp Leu Pro Ala Pro Val Leu Glu Asn
 165 170 175

Val Gly Arg Phe Arg Lys Lys Ala Glu Gly Leu Thr Gly Trp Arg Phe
 180 185 190

Ser Gly Gly Ile Ser Pro Ala Val Asn Arg Asn Ala Asn Asn Ala Ala
 195 200 205

Pro Gln Tyr Cys Arg Gln Asn Gly Gly Arg Gln Ile Cys Ser Val Ser
 210 215 220

Arg Ala Glu Arg Ala Ala Gly Leu Asn Tyr Glu Ile Glu Ala Glu Lys
 225 230 235 240

Leu Thr Ala Leu Ala Asp Asn His Tyr Leu Leu Phe Arg Ser Asn Ile
 245 250 255

Gly Gly Thr Ser Tyr Tyr Phe Ser Lys Lys Ser Ala Tyr Asp Asp Gly
 260 265 270

Phe Gly Arg Ala Tyr Leu Gly Trp Gln Tyr Lys Asn Ala Arg Gln Thr
 275 280 285

Ala Gly Ile Leu Pro Phe Tyr Gln Val Gln Leu Ser Gly Ser Asp Gly
 290 295 300

Phe Asp Ala Lys Thr Lys Arg Val Asn Asn Arg Arg Leu Pro Pro Tyr
305 310 315 320

Met Leu Ala His Gly Val Gly Val Gln Leu Ser His Thr Tyr Arg Pro
325 330 335

Asn Pro Gly Trp Gln Phe Ser Val Ala Leu Glu His Tyr Arg Gln Arg
340 345 350

Tyr Arg Glu Gln Asp Arg Ala Glu Tyr Asn Asn Gly Arg Gln Asp Gly
355 360 365

Phe Tyr Val Ser Ser Ala Lys Arg Leu Gly Glu Ser Ala Thr Val Phe
370 375 380

Gly Gly Trp Gln Phe Val Arg Phe Val Pro Lys Arg Glu Thr Val Gly
385 390 395 400

Gly Ala Val Asn Asn Ala Ala Tyr Arg Arg Asn Gly Val Tyr Ala Gly
405 410 415

Trp Ala Gln Glu Trp Arg Gln Leu Gly Gly Leu Asn Ser Arg Val Ser
420 425 430

Ala Ser Tyr Ala Arg Arg Asn Tyr Lys Gly Val Ala Ala Phe Ser Thr
435 440 445

Glu Ala Gln Arg Asn Arg Glu Trp Asn Val Ser Leu Ala Leu Ser His
450 455 460

Asp Lys Leu Ser Tyr Lys Gly Ile Val Pro Ala Leu Asn Tyr Arg Phe
465 470 475 480

Gly Arg Thr Glu Ser Asn Val Pro Tyr Ala Lys Arg Arg Asn Ser Glu
485 490 495

Val Phe Val Ser Ala Asp Trp Arg Phe
500 505

<210> 145

<211> 840

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(840)

<400> 145

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Met Asp Lys Glu Arg Ile Leu Thr Pro Ala Val Val Phe Ser Val Ala	
1 5 10 15	
ctg ctg cat ttg gca atg gtg gca ttg ctc tgg cag gcg cac aag ctg	96
Leu Leu His Leu Ala Met Val Ala Leu Leu Trp Gln Ala His Lys Leu	
20 25 30	
ccc gtg ata gag tca ggc aat gtt att gaa ttt gtc gat ttg ggc gat	144
Pro Val Ile Glu Ser Gly Asn Val Ile Glu Phe Val Asp Leu Gly Asp	
35 40 45	
ttt ggc gga ggg gac ggc gca ccc gaa ggt gca ggc gcg cct gcc gcg	192
Phe Gly Gly Gly Asp Gly Ala Pro Glu Gly Ala Gly Ala Pro Ala Ala	
50 55 60	
ccc gaa ccg caa ccc gtg ccc gag ccg ccc aaa cct gtc gag ccg ccc	240
Pro Glu Pro Gln Pro Val Pro Glu Pro Pro Lys Pro Val Glu Pro Pro	
65 70 75 80	
aag ccg gtt ttg aag ccg gtg gtt acg aaa aag gcg gat gcg gat att	288
Lys Pro Val Leu Lys Pro Val Val Thr Lys Lys Ala Asp Ala Asp Ile	
85 90 95	
cag cag cct aag gaa gag ccg aaa cct gaa gaa aag ccg aaa ccc gaa	336
Gln Gln Pro Lys Glu Glu Pro Lys Pro Glu Glu Lys Pro Lys Pro Glu	
100 105 110	
gaa aaa ccg aaa cca gag cct aag ccg gaa gcg aag cct gtc ccg aaa	384
Glu Lys Pro Lys Pro Glu Pro Lys Pro Glu Ala Lys Pro Val Pro Lys	
115 120 125	
ccg gcg gaa aaa ccg gtc gag aag ccg tct gaa aaa cct gcc gaa cat	432
Pro Ala Glu Lys Pro Val Glu Lys Pro Ser Glu Lys Pro Ala Glu His	
130 135 140	
ccc ggc aat gct tct gcc aaa gca gac agc gag cag ggc aat ggg gaa	480
Pro Gly Asn Ala Ser Ala Lys Ala Asp Ser Glu Gln Gly Asn Gly Glu	
145 150 155 160	
gat aag gga acc ggt atc aaa gga gac gga acg ggg cgc gga gaa ggc	528
Asp Lys Gly Thr Gly Ile Lys Gly Asp Gly Thr Gly Arg Gly Glu Gly	
165 170 175	
agc ggt aaa ggt agc ggc ggt gtc aaa ggc gaa cac ggg gaa gga gcc	576

Ser Gly Lys Gly Ser Gly Gly Val Lys Gly Glu His Gly Glu Gly Ala
 180 185 190
 ggc agc agc aaa ggc aat cct tta cgc gcc aac ggc agc att ccg cgc 624
 Gly Ser Ser Lys Gly Asn Pro Leu Arg Ala Asn Gly Ser Ile Pro Arg
 195 200 205
 ccg gct tat ccc acg ctt tct atg gag aat gat gag cag ggt acg gtt 672
 Pro Ala Tyr Pro Thr Leu Ser Met Glu Asn Asp Glu Gln Gly Thr Val
 210 215 220
 gtt ttg agc gtt ttg gtt tct ccg ggc ggt cat gtt gag tcc gtt aaa 720
 Val Leu Ser Val Leu Val Ser Pro Gly Gly His Val Glu Ser Val Lys
 225 230 235 240
 atc gtg aaa agc agt ggt ttt tcc cgg ttg gac aat gcg gca cgc aag 768
 Ile Val Lys Ser Ser Gly Phe Ser Arg Leu Asp Asn Ala Ala Arg Lys
 245 250 255
 gcg gcg caa aac ggg cat ttt caa gcc aat gcc tgg acg gag ttt aaa 816
 Ala Ala Gln Asn Gly His Phe Gln Ala Asn Ala Trp Thr Glu Phe Lys
 260 265 270
 gtc ccc gtc aag ttt gaa ttg aat 840
 Val Pro Val Lys Phe Glu Leu Asn
 275 280
 <210> 146
 <211> 280
 <212> PRT
 <213> Neisseria meningitidis
 <400> 146
 Met Asp Lys Glu Arg Ile Leu Thr Pro Ala Val Val Phe Ser Val Ala
 1 5 10 15
 Leu Leu His Leu Ala Met Val Ala Leu Leu Trp Gln Ala His Lys Leu
 20 25 30
 Pro Val Ile Glu Ser Gly Asn Val Ile Glu Phe Val Asp Leu Gly Asp
 35 40 45
 Phe Gly Gly Gly Asp Gly Ala Pro Glu Gly Ala Gly Ala Pro Ala Ala
 50 55 60
 Pro Glu Pro Gln Pro Val Pro Glu Pro Pro Lys Pro Val Glu Pro Pro
 65 70 75 80

Lys Pro Val Leu Lys Pro Val Val Thr Lys Lys Ala Asp Ala Asp Ile
85 90 95

Gln Gln Pro Lys Glu Glu Pro Lys Pro Glu Glu Lys Pro Lys Pro Glu
100 105 110

Glu Lys Pro Lys Pro Glu Pro Lys Pro Glu Ala Lys Pro Val Pro Lys
115 120 125

Pro Ala Glu Lys Pro Val Glu Lys Pro Ser Glu Lys Pro Ala Glu His
130 135 140

Pro Gly Asn Ala Ser Ala Lys Ala Asp Ser Glu Gln Gly Asn Gly Glu
145 150 155 160

Asp Lys Gly Thr Gly Ile Lys Gly Asp Gly Thr Gly Arg Gly Glu Gly
165 170 175

Ser Gly Lys Gly Ser Gly Gly Val Lys Gly Glu His Gly Glu Gly Ala
180 185 190

Gly Ser Ser Lys Gly Asn Pro Leu Arg Ala Asn Gly Ser Ile Pro Arg
195 200 205

Pro Ala Tyr Pro Thr Leu Ser Met Glu Asn Asp Glu Gln Gly Thr Val
210 215 220

Val Leu Ser Val Leu Val Ser Pro Gly Gly His Val Glu Ser Val Lys
225 230 235 240

Ile Val Lys Ser Ser Gly Phe Ser Arg Leu Asp Asn Ala Ala Arg Lys
245 250 255

Ala Ala Gln Asn Gly His Phe Gln Ala Asn Ala Trp Thr Glu Phe Lys
260 265 270

Val Pro Val Lys Phe Glu Leu Asn
275 280

<210> 147

<211> 1572

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1572)

<400> 147

atg ttc aga cgg tat ctt ccg aac aga cag atg aat atg gtt tcc aaa	48
Met Phe Arg Arg Tyr Leu Pro Asn Arg Gln Met Asn Met Val Ser Lys	
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ctg gac aaa tac tgg cag cac ccc gcc ctc tac tgg cct ttg ctc atc	96
Leu Asp Lys Tyr Trp Gln His Pro Ala Leu Tyr Trp Pro Leu Leu Ile	
20 25 30	
ctt ttt gcc gcc gcc acc ccc ttt acc ttc gca ccc tac tac cac ttt	144
Leu Phe Ala Ala Ala Thr Pro Phe Thr Phe Ala Pro Tyr Tyr His Phe	
35 40 45	
tgg ctg atg ccc ttg att ttc ggt gcc ttc gtc cgc ctc atc gaa ctg	192
Trp Leu Met Pro Leu Ile Phe Gly Ala Phe Val Arg Leu Ile Glu Leu	
50 55 60	
cgt ccg cgt ttt gct gtc tct tcc gcc tac ctg ttc ggc ctg acc gca	240
Arg Pro Arg Phe Ala Val Ser Ser Ala Tyr Leu Phe Gly Leu Thr Ala	
65 70 75 80	
tac acg aca cag ttc tac tgg ata cac acc gcc ctg cac gac gtt tcc	288
Tyr Thr Thr Gln Phe Tyr Trp Ile His Thr Ala Leu His Asp Val Ser	
85 90 95	
ggc ctg ccc gac ctc tat gcc gta ccg ctg acc ttc cta ctc ccc gcc	336
Gly Leu Pro Asp Leu Tyr Ala Val Pro Leu Thr Phe Leu Leu Pro Ala	
100 105 110	
tac ctt gcc ctt tat ccg gca ctg tgt ttc tgg ctg tgg aaa aaa ttt	384
Tyr Leu Ala Leu Tyr Pro Ala Leu Cys Phe Trp Leu Trp Lys Lys Phe	
115 120 125	
acc ctg cct cgg ggc ata aaa atc ggt ttg gta ctg ccc atc ctg tgg	432
Thr Leu Pro Arg Gly Ile Lys Ile Gly Leu Val Leu Pro Ile Leu Trp	
130 135 140	
aca ctg acc gag ttt gcc cgc gaa cgt ttc ctg acc gga ttc ggc tgg	480
Thr Leu Thr Glu Phe Ala Arg Glu Arg Phe Leu Thr Gly Phe Gly Trp	
145 150 155 160	
ggc gca atc ggc tac tcc caa atc acc ccg gac agc ccg ctc gcc ggc	528
Gly Ala Ile Gly Tyr Ser Gln Ile Thr Pro Asp Ser Pro Leu Ala Gly	
165 170 175	

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ttt gcc cca ttt ggc ggc atc cac atg gtt aca ctg gca acc gcc ttt 576
Phe Ala Pro Phe Gly Gly Ile His Met Val Thr Leu Ala Thr Ala Phe
      180              185              190

ctc ggt gtc tgg ctg gtt ttg gcg agt gat aac acc gca cgt tcg ggc 624
Leu Gly Val Trp Leu Val Leu Ala Ser Asp Asn Thr Ala Arg Ser Gly
      195              200              205

aaa cgc ctg ctt cca att atc ctg att gcc gcc ctg ctt gcc gca ggc 672
Lys Arg Leu Leu Pro Ile Ile Leu Ile Ala Ala Leu Leu Ala Ala Gly
      210              215              220

tac acc gcc cga caa acc gac ttc acc cgc ccc gac ggc agc cgc agc 720
Tyr Thr Ala Arg Gln Thr Asp Phe Thr Arg Pro Asp Gly Ser Arg Ser
      225              230              235              240

acc gtc gcc ctg ctt caa ggc aac atc gac caa acc ctc aaa tgg cgt 768
Thr Val Ala Leu Leu Gln Gly Asn Ile Asp Gln Thr Leu Lys Trp Arg
      245              250              255

gaa gac caa gtc atc ccg acc ata cag aaa tat tac gaa cag gtc ggc 816
Glu Asp Gln Val Ile Pro Thr Ile Gln Lys Tyr Tyr Glu Gln Val Gly
      260              265              270

aaa acc act gcc gac atc gtc atc ctg ccc gaa acc gcc atc ccc gtc 864
Lys Thr Thr Ala Asp Ile Val Ile Leu Pro Glu Thr Ala Ile Pro Val
      275              280              285

atg cgc caa aac ctg ccg gaa aac ata ctg gca aaa ttt gcc gaa cag 912
Met Arg Gln Asn Leu Pro Glu Asn Ile Leu Ala Lys Phe Ala Glu Gln
      290              295              300

gcg caa aac aac ggc agc gcg ctc gcc gtc ggc atc agc caa tac act 960
Ala Gln Asn Asn Gly Ser Ala Leu Ala Val Gly Ile Ser Gln Tyr Thr
      305              310              315              320

tcg gac ggc aac ggt tac gaa aac gcc gtc atc aac ctg acc ggt tat 1008
Ser Asp Gly Asn Gly Tyr Glu Asn Ala Val Ile Asn Leu Thr Gly Tyr
      325              330              335

cag gaa aac aat cag gac ggc atc ccc tac tac gcc aaa aac cac ctc 1056
Gln Glu Asn Asn Gln Asp Gly Ile Pro Tyr Tyr Ala Lys Asn His Leu
      340              345              350

gtc ccc ttc ggc gaa tac aaa ccg ctg ccc ttc ctg acc acg ccg ctt 1104
Val Pro Phe Gly Glu Tyr Lys Pro Leu Pro Phe Leu Thr Thr Pro Leu
      355              360              365

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tac aaa atg atg aat atg ccc ctt tcc gac ttc cgc aaa ggc ggc ggt 1152
 Tyr Lys Met Met Asn Met Pro Leu Ser Asp Phe Arg Lys Gly Gly Gly
 370 375 380

aag caa tcc gcc ctg ctg atg aaa aac caa aaa atc gcc ttc aac atc 1200
 Lys Gln Ser Ala Leu Leu Met Lys Asn Gln Lys Ile Ala Phe Asn Ile
 385 390 395 400

tgt tac gaa gac gga ttc ggc gac gaa ctg att gcc gcc gcc aaa gat 1248
 Cys Tyr Glu Asp Gly Phe Gly Asp Glu Leu Ile Ala Ala Ala Lys Asp
 405 410 415

gcc aca ctg ctt gcc aat gcc agc aat atg gcg tgg tac gga aaa tcc 1296
 Ala Thr Leu Leu Ala Asn Ala Ser Asn Met Ala Trp Tyr Gly Lys Ser
 420 425 430

aac gcc atg tac cag cac ctc caa caa tcg cag gcg cgg gct atg gaa 1344
 Asn Ala Met Tyr Gln His Leu Gln Gln Ser Gln Ala Arg Ala Met Glu
 435 440 445

ctc gga cgc tat atg gtc cgt gcc acc aac acc ggc gca acc gcc atc 1392
 Leu Gly Arg Tyr Met Val Arg Ala Thr Asn Thr Gly Ala Thr Ala Ile
 450 455 460

atc tcc ccc aaa ggc aac atc atc gcc caa gcc caa ccc gat acg gaa 1440
 Ile Ser Pro Lys Gly Asn Ile Ile Ala Gln Ala Gln Pro Asp Thr Glu
 465 470 475 480

acc gta ttg gaa gga cac atc aaa ggc tat gtc ggc gaa acg ccc tat 1488
 Thr Val Leu Glu Gly His Ile Lys Gly Tyr Val Gly Glu Thr Pro Tyr
 485 490 495

atg aaa acc ggc agt tca tgg tgg ttg atg ggc ata ttg acc cta gcc 1536
 Met Lys Thr Gly Ser Ser Trp Trp Leu Met Gly Ile Leu Thr Leu Ala
 500 505 510

gca ctg att ctt ttc atc ttc cga aac aaa gaa cac 1572
 Ala Leu Ile Leu Phe Ile Phe Arg Asn Lys Glu His
 515 520

<210> 148

<211> 524

<212> PRT

<213> Neisseria meningitidis

<400> 148

Met Phe Arg Arg Tyr Leu Pro Asn Arg Gln Met Asn Met Val Ser Lys

1	5	10	15
Leu Asp Lys Tyr Trp Gln His Pro Ala Leu Tyr Trp Pro Leu Leu Ile	20	25	30
Leu Phe Ala Ala Ala Thr Pro Phe Thr Phe Ala Pro Tyr Tyr His Phe	35	40	45
Trp Leu Met Pro Leu Ile Phe Gly Ala Phe Val Arg Leu Ile Glu Leu	50	55	60
Arg Pro Arg Phe Ala Val Ser Ser Ala Tyr Leu Phe Gly Leu Thr Ala	65	70	75
Tyr Thr Thr Gln Phe Tyr Trp Ile His Thr Ala Leu His Asp Val Ser	85	90	95
Gly Leu Pro Asp Leu Tyr Ala Val Pro Leu Thr Phe Leu Leu Pro Ala	100	105	110
Tyr Leu Ala Leu Tyr Pro Ala Leu Cys Phe Trp Leu Trp Lys Lys Phe	115	120	125
Thr Leu Pro Arg Gly Ile Lys Ile Gly Leu Val Leu Pro Ile Leu Trp	130	135	140
Thr Leu Thr Glu Phe Ala Arg Glu Arg Phe Leu Thr Gly Phe Gly Trp	145	150	155
Gly Ala Ile Gly Tyr Ser Gln Ile Thr Pro Asp Ser Pro Leu Ala Gly	165	170	175
Phe Ala Pro Phe Gly Gly Ile His Met Val Thr Leu Ala Thr Ala Phe	180	185	190
Leu Gly Val Trp Leu Val Leu Ala Ser Asp Asn Thr Ala Arg Ser Gly	195	200	205
Lys Arg Leu Leu Pro Ile Ile Leu Ile Ala Ala Leu Leu Ala Ala Gly	210	215	220
Tyr Thr Ala Arg Gln Thr Asp Phe Thr Arg Pro Asp Gly Ser Arg Ser	225	230	235
Thr Val Ala Leu Leu Gln Gly Asn Ile Asp Gln Thr Leu Lys Trp Arg	245	250	255
Glu Asp Gln Val Ile Pro Thr Ile Gln Lys Tyr Tyr Glu Gln Val Gly			

260	265	270
Lys Thr Thr Ala Asp Ile Val Ile Leu Pro Glu Thr Ala Ile Pro Val		
275	280	285
Met Arg Gln Asn Leu Pro Glu Asn Ile Leu Ala Lys Phe Ala Glu Gln		
290	295	300
Ala Gln Asn Asn Gly Ser Ala Leu Ala Val Gly Ile Ser Gln Tyr Thr		
305	310	315
Ser Asp Gly Asn Gly Tyr Glu Asn Ala Val Ile Asn Leu Thr Gly Tyr		
325	330	335
Gln Glu Asn Asn Gln Asp Gly Ile Pro Tyr Tyr Ala Lys Asn His Leu		
340	345	350
Val Pro Phe Gly Glu Tyr Lys Pro Leu Pro Phe Leu Thr Thr Pro Leu		
355	360	365
Tyr Lys Met Met Asn Met Pro Leu Ser Asp Phe Arg Lys Gly Gly Gly		
370	375	380
Lys Gln Ser Ala Leu Leu Met Lys Asn Gln Lys Ile Ala Phe Asn Ile		
385	390	395
Cys Tyr Glu Asp Gly Phe Gly Asp Glu Leu Ile Ala Ala Ala Lys Asp		
405	410	415
Ala Thr Leu Leu Ala Asn Ala Ser Asn Met Ala Trp Tyr Gly Lys Ser		
420	425	430
Asn Ala Met Tyr Gln His Leu Gln Gln Ser Gln Ala Arg Ala Met Glu		
435	440	445
Leu Gly Arg Tyr Met Val Arg Ala Thr Asn Thr Gly Ala Thr Ala Ile		
450	455	460
Ile Ser Pro Lys Gly Asn Ile Ile Ala Gln Ala Gln Pro Asp Thr Glu		
465	470	475
Thr Val Leu Glu Gly His Ile Lys Gly Tyr Val Gly Glu Thr Pro Tyr		
485	490	495
Met Lys Thr Gly Ser Ser Trp Trp Leu Met Gly Ile Leu Thr Leu Ala		
500	505	510
Ala Leu Ile Leu Phe Ile Phe Arg Asn Lys Glu His		

515

520

<210> 149

<211> 558

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(558)

<400> 149

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ttg gat ctc tac aaa ctc ggt tac aaa caa cat acc cgt acc gat gcg      48
Leu Asp Leu Tyr Lys Leu Gly Tyr Lys Gln His Thr Arg Thr Asp Ala
  1              5              10              15

gaa ggc tat atc gag aaa ctg cac att acc ccc gcc aat gcc cat gag      96
Glu Gly Tyr Ile Glu Lys Leu His Ile Thr Pro Ala Asn Ala His Glu
              20              25              30

tgc aaa cac ctg ccg ccg ttg ttg gaa gga ctg ccc aaa ggt acg acc      144
Cys Lys His Leu Pro Pro Leu Leu Glu Gly Leu Pro Lys Gly Thr Thr
              35              40              45

gtc tat gcc gac aaa ggc tac gac agt gcg gaa aac cgg caa cat ctg      192
Val Tyr Ala Asp Lys Gly Tyr Asp Ser Ala Glu Asn Arg Gln His Leu
              50              55              60

gaa gaa cat cag ttg ttg gac ggc att atg cgc aaa gcc tgc cgc aac      240
Glu Glu His Gln Leu Leu Asp Gly Ile Met Arg Lys Ala Cys Arg Asn
              65              70              75              80

cgt ccg ctg tcg gaa acg caa acc aaa cgc aac cgg tat ttg tcg aag      288
Arg Pro Leu Ser Glu Thr Gln Thr Lys Arg Asn Arg Tyr Leu Ser Lys
              85              90              95

acc cgt tat agt gga tta aat tta aat cag gac aag gcg acg aag ccg      336
Thr Arg Tyr Ser Gly Leu Asn Leu Asn Gln Asp Lys Ala Thr Lys Pro
              100              105              110

cag aca gta caa ata gta cgg caa ggc gag gca acg ccg tac tgg ttt      384
Gln Thr Val Gln Ile Val Arg Gln Gly Glu Ala Thr Pro Tyr Trp Phe
              115              120              125

aaa ttt aat cca cta tat gtg gtc gaa cag agc ttc ggt acg ctg cac      432
Lys Phe Asn Pro Leu Tyr Val Val Glu Gln Ser Phe Gly Thr Leu His

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130 135 140
 cgt aaa ttc cgc tat gcg cgg gca gcc tat ttc gga ctg att aaa gtg 480
 Arg Lys Phe Arg Tyr Ala Arg Ala Ala Tyr Phe Gly Leu Ile Lys Val
 145 150 155 160
 agt gcg caa agc cat ctg aag gcg atg tgt ttg aac ctg ttg aaa gcc 528
 Ser Ala Gln Ser His Leu Lys Ala Met Cys Leu Asn Leu Leu Lys Ala
 165 170 175
 gcc aac aag cta agt gcg ccc gct gcc gcc 558
 Ala Asn Lys Leu Ser Ala Pro Ala Ala Ala
 180 185

 <210> 150
 <211> 186
 <212> PRT
 <213> Neisseria meningitidis

 <400> 150
 Leu Asp Leu Tyr Lys Leu Gly Tyr Lys Gln His Thr Arg Thr Asp Ala
 1 5 10 15
 Glu Gly Tyr Ile Glu Lys Leu His Ile Thr Pro Ala Asn Ala His Glu
 20 25 30
 Cys Lys His Leu Pro Pro Leu Leu Glu Gly Leu Pro Lys Gly Thr Thr
 35 40 45
 Val Tyr Ala Asp Lys Gly Tyr Asp Ser Ala Glu Asn Arg Gln His Leu
 50 55 60
 Glu Glu His Gln Leu Leu Asp Gly Ile Met Arg Lys Ala Cys Arg Asn
 65 70 75 80
 Arg Pro Leu Ser Glu Thr Gln Thr Lys Arg Asn Arg Tyr Leu Ser Lys
 85 90 95
 Thr Arg Tyr Ser Gly Leu Asn Leu Asn Gln Asp Lys Ala Thr Lys Pro
 100 105 110
 Gln Thr Val Gln Ile Val Arg Gln Gly Glu Ala Thr Pro Tyr Trp Phe
 115 120 125
 Lys Phe Asn Pro Leu Tyr Val Val Glu Gln Ser Phe Gly Thr Leu His
 130 135 140

Arg Lys Phe Arg Tyr Ala Arg Ala Ala Tyr Phe Gly Leu Ile Lys Val
 145 150 155 160

Ser Ala Gln Ser His Leu Lys Ala Met Cys Leu Asn Leu Leu Lys Ala
 165 170 175

Ala Asn Lys Leu Ser Ala Pro Ala Ala Ala
 180 185

<210> 151

<211> 1332

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1332)

<400> 151

atg act gac ctg aac acc ctg ttt gcc aac ctc aaa caa cgc aat ccc 48
 Met Thr Asp Leu Asn Thr Leu Phe Ala Asn Leu Lys Gln Arg Asn Pro
 1 5 10 15

aat cag gag ccg ttc cat cag gcg gtt gaa gaa gtc ttc atg agt ctc 96
 Asn Gln Glu Pro Phe His Gln Ala Val Glu Glu Val Phe Met Ser Leu
 20 25 30

gat ccg ttt ttg gca aaa aat ccg aaa tac acc cag caa agc ctg ctg 144
 Asp Pro Phe Leu Ala Lys Asn Pro Lys Tyr Thr Gln Gln Ser Leu Leu
 35 40 45

gaa cgc atc gtc gaa ccc gaa cgc gtc gtg atg ttc cgc gta acc tgg 192
 Glu Arg Ile Val Glu Pro Glu Arg Val Val Met Phe Arg Val Thr Trp
 50 55 60

cag gac gat aaa ggg caa gtc caa gtc aac cgg ggc tac cgc gtg caa 240
 Gln Asp Asp Lys Gly Gln Val Gln Val Asn Arg Gly Tyr Arg Val Gln
 65 70 75 80

atg agt tcc gcc atc ggt cct tac aaa ggc ggc ctg cgc ttc cat ccg 288
 Met Ser Ser Ala Ile Gly Pro Tyr Lys Gly Gly Leu Arg Phe His Pro
 85 90 95

acc gtc gat ttg ggc gta ttg aaa ttc ctc gct ttt gaa caa gtg ttc 336
 Thr Val Asp Leu Gly Val Leu Lys Phe Leu Ala Phe Glu Gln Val Phe
 100 105 110

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aaa aac gcc ttg acc acc ctg cct atg ggc ggc ggc aaa ggc ggt tcc 384
Lys Asn Ala Leu Thr Thr Leu Pro Met Gly Gly Gly Lys Gly Gly Ser
      115                120                125

gac ttc gac ccc aaa ggc aaa tcc gat gcc gaa gta atg cgc ttc tgc 432
Asp Phe Asp Pro Lys Gly Lys Ser Asp Ala Glu Val Met Arg Phe Cys
      130                135                140

caa gcc ttt atg acc gaa ctc tac cgc cac atc ggc gcg gac acc gat 480
Gln Ala Phe Met Thr Glu Leu Tyr Arg His Ile Gly Ala Asp Thr Asp
      145                150                155                160

gtt ccg gcc ggc gac atc ggc gta ggc ggg cgc gaa atc ggc tac ctg 528
Val Pro Ala Gly Asp Ile Gly Val Gly Gly Arg Glu Ile Gly Tyr Leu
      165                170                175

ttc gga caa tac aaa aaa atc cgc aac gag ttt tct tcc gtc ctg acc 576
Phe Gly Gln Tyr Lys Lys Ile Arg Asn Glu Phe Ser Ser Val Leu Thr
      180                185                190

ggc aaa ggt ttg gaa tgg ggc ggc agc ctc atc cgt ccc gaa gcg acc 624
Gly Lys Gly Leu Glu Trp Gly Gly Ser Leu Ile Arg Pro Glu Ala Thr
      195                200                205

ggc tac ggc tgc gtc tat ttc gcc caa gcg atg ctg caa acc cgc aac 672
Gly Tyr Gly Cys Val Tyr Phe Ala Gln Ala Met Leu Gln Thr Arg Asn
      210                215                220

gat agt ttt gaa ggc aaa cgc gtc ctg att tcc ggt tcc ggc aac gtg 720
Asp Ser Phe Glu Gly Lys Arg Val Leu Ile Ser Gly Ser Gly Asn Val
      225                230                235                240

gcg caa tac gcc gcc gaa aaa gcc atc caa ctg ggc gcg aaa gtg ctg 768
Ala Gln Tyr Ala Ala Glu Lys Ala Ile Gln Leu Gly Ala Lys Val Leu
      245                250                255

acc gtt tcc gac tcc aac ggc ttc gtc ctc ttc ccc gac agc ggt atg 816
Thr Val Ser Asp Ser Asn Gly Phe Val Leu Phe Pro Asp Ser Gly Met
      260                265                270

agc gaa gcg caa ctc gcc gcc ttg atc gaa ttg aaa gaa gtc cgc cgc 864
Ser Glu Ala Ala Gln Leu Ala Ala Leu Ile Glu Leu Lys Glu Val Arg Arg
      275                280                285

gaa cgg gtt gcc acc tac gcc aaa gag caa ggt ctg caa tac ttt gaa 912
Glu Arg Val Ala Thr Tyr Ala Lys Glu Gln Gly Leu Gln Tyr Phe Glu
      290                295                300

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aac caa aaa ccg tgg ggc gtc gcc gcc gaa atc gcc ctg ccc tgc gcg 960
 Asn Gln Lys Pro Trp Gly Val Ala Ala Glu Ile Ala Leu Pro Cys Ala
 305 310 315 320

 acc cag aac gaa ttg gac gaa gaa gcc gcc aaa acc ctg ttg gca aac 1008
 Thr Gln Asn Glu Leu Asp Glu Glu Ala Ala Lys Thr Leu Leu Ala Asn
 325 330 335

 ggc tgc tac gtc gtt gcc gaa ggt gcg aat atg ccg tcg act ttg ggc 1056
 Gly Cys Tyr Val Val Ala Glu Gly Ala Asn Met Pro Ser Thr Leu Gly
 340 345 350

 gcg gtc gag caa ttt atc aaa gcc ggc atc ctc tac gcc ccg ggc aaa 1104
 Ala Val Glu Gln Phe Ile Lys Ala Gly Ile Leu Tyr Ala Pro Gly Lys
 355 360 365

 gcc tcc aac gcc ggc ggc gta gca act tca ggt ttg gaa atg agc caa 1152
 Ala Ser Asn Ala Gly Gly Val Ala Thr Ser Gly Leu Glu Met Ser Gln
 370 375 380

 aac gcc atc cgc ctg tct tgg act cgt gaa gaa gtc gac caa cgc ctg 1200
 Asn Ala Ile Arg Leu Ser Trp Thr Arg Glu Glu Val Asp Gln Arg Leu
 385 390 395 400

 ttc ggc atc atg caa agc atc cac gaa tcc tgt ctg aaa tac ggc aaa 1248
 Phe Gly Ile Met Gln Ser Ile His Glu Ser Cys Leu Lys Tyr Gly Lys
 405 410 415

 gtc ggc gac aca gta aac tac gtc aat ggt gcg aac att gcc ggt ttc 1296
 Val Gly Asp Thr Val Asn Tyr Val Asn Gly Ala Asn Ile Ala Gly Phe
 420 425 430

 gtc aaa gtt gcc gat gcg atg ctg gcg caa ggc ttc 1332
 Val Lys Val Ala Asp Ala Met Leu Ala Gln Gly Phe
 435 440

<210> 152

<211> 444

<212> PRT

<213> Neisseria meningitidis

<400> 152

Met Thr Asp Leu Asn Thr Leu Phe Ala Asn Leu Lys Gln Arg Asn Pro
 1 5 10 15

Asn Gln Glu Pro Phe His Gln Ala Val Glu Glu Val Phe Met Ser Leu

302

275 280 285
 Glu Arg Val Ala Thr Tyr Ala Lys Glu Gln Gly Leu Gln Tyr Phe Glu
 290 295 300
 Asn Gln Lys Pro Trp Gly Val Ala Ala Glu Ile Ala Leu Pro Cys Ala
 305 310 315 320
 Thr Gln Asn Glu Leu Asp Glu Glu Ala Ala Lys Thr Leu Leu Ala Asn
 325 330 335
 Gly Cys Tyr Val Val Ala Glu Gly Ala Asn Met Pro Ser Thr Leu Gly
 340 345 350
 Ala Val Glu Gln Phe Ile Lys Ala Gly Ile Leu Tyr Ala Pro Gly Lys
 355 360 365
 Ala Ser Asn Ala Gly Gly Val Ala Thr Ser Gly Leu Glu Met Ser Gln
 370 375 380
 Asn Ala Ile Arg Leu Ser Trp Thr Arg Glu Glu Val Asp Gln Arg Leu
 385 390 395 400
 Phe Gly Ile Met Gln Ser Ile His Glu Ser Cys Leu Lys Tyr Gly Lys
 405 410 415
 Val Gly Asp Thr Val Asn Tyr Val Asn Gly Ala Asn Ile Ala Gly Phe
 420 425 430
 Val Lys Val Ala Asp Ala Met Leu Ala Gln Gly Phe
 435 440

<210> 153

<211> 867

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(867)

<400> 153

atg aaa ccc ata cgg aaa gcc gtc ttc ccc gtc gca ggg atg gga aca 48
 Met Lys Pro Ile Arg Lys Ala Val Phe Pro Val Ala Gly Met Gly Thr
 1 5 10 15

cgc ttc ctg ccc gcc acc aag gca agc ccg aaa gaa atg ctg ccc atc	96
Arg Phe Leu Pro Ala Thr Lys Ala Ser Pro Lys Glu Met Leu Pro Ile	
20 25 30	
gtc gac aag ccg ctg atc caa tac gcc gta gaa gaa gcc gtg gaa gcc	144
Val Asp Lys Pro Leu Ile Gln Tyr Ala Val Glu Glu Ala Val Glu Ala	
35 40 45	
ggc tgc acg gaa atg gtg ttt gtt acc gga cgc aac aaa cgc agc atc	192
Gly Cys Thr Glu Met Val Phe Val Thr Gly Arg Asn Lys Arg Ser Ile	
50 55 60	
gaa gac cat ttc gac aag gca tac gaa ctc gaa acc gag ttg gaa atg	240
Glu Asp His Phe Asp Lys Ala Tyr Glu Leu Glu Thr Glu Leu Glu Met	
65 70 75 80	
cgc cat aaa gac aaa ttg ttg gaa cac gtc cgc aac atc ctg ccg ccg	288
Arg His Lys Asp Lys Leu Leu Glu His Val Arg Asn Ile Leu Pro Pro	
85 90 95	
aac att acc tgc ctc tac atc cgt cag gcg gaa gca ctg ggc ttg gga	336
Asn Ile Thr Cys Leu Tyr Ile Arg Gln Ala Glu Ala Leu Gly Leu Gly	
100 105 110	
cac gcc gtc ttg tgc gcc cgc gcc gcc atc ggc gac gaa ccc ttt gcc	384
His Ala Val Leu Cys Ala Arg Ala Ala Ile Gly Asp Glu Pro Phe Ala	
115 120 125	
gtg att ttg gct gac gat tta atc gat gcc caa aaa ggc gcg ctc aaa	432
Val Ile Leu Ala Asp Asp Leu Ile Asp Ala Gln Lys Gly Ala Leu Lys	
130 135 140	
caa atg gtc gaa gtg tac gaa cgc agc ggc aac agc att ttg ggc gta	480
Gln Met Val Glu Val Tyr Glu Arg Ser Gly Asn Ser Ile Leu Gly Val	
145 150 155 160	
gaa act gtc gaa ccg tcg caa acc ggc tca tac ggc atc gtc gaa acc	528
Glu Thr Val Glu Pro Ser Gln Thr Gly Ser Tyr Gly Ile Val Glu Thr	
165 170 175	
gaa cag ctc aaa cag ttc caa cgc att acc ggc att gtc gaa aaa ccc	576
Glu Gln Leu Lys Gln Phe Gln Arg Ile Thr Gly Ile Val Glu Lys Pro	
180 185 190	
aag ccc gaa gac gcg ccc tcc aac ctt gcc gtc gtc gga cgc tac atc	624
Lys Pro Glu Asp Ala Pro Ser Asn Leu Ala Val Val Gly Arg Tyr Ile	
195 200 205	

ctt acc ccg cgc att ttc gac tta ctg acc gga ctg ccg cgc ggc gcg 672
 Leu Thr Pro Arg Ile Phe Asp Leu Leu Thr Gly Leu Pro Arg Gly Ala
 210 215 220

ggc aac gaa atc cag ctt aca gac ggc atc gcc aag ctg ctc gat cac 720
 Gly Asn Glu Ile Gln Leu Thr Asp Gly Ile Ala Lys Leu Leu Asp His
 225 230 235 240

gaa ttt gtc cta gcg cac ccc ttt gaa ggc acg cgc tac gac tgc ggc 768
 Glu Phe Val Leu Ala His Pro Phe Glu Gly Thr Arg Tyr Asp Cys Gly
 245 250 255

agc aaa ctg ggc tat ctg gaa gcc acc gtc gcc tac ggc ctg aaa cac 816
 Ser Lys Leu Gly Tyr Leu Glu Ala Thr Val Ala Tyr Gly Leu Lys His
 260 265 270

ccc gaa acc ggc gaa ccc ttc cgc cgc ctt ttg gaa aaa tac cgt acc 864
 Pro Glu Thr Gly Glu Pro Phe Arg Arg Leu Leu Glu Lys Tyr Arg Thr
 275 280 285

gaa 867
 Glu

<210> 154

<211> 289

<212> PRT

<213> Neisseria meningitidis

<400> 154

Met Lys Pro Ile Arg Lys Ala Val Phe Pro Val Ala Gly Met Gly Thr
 1 5 10 15

Arg Phe Leu Pro Ala Thr Lys Ala Ser Pro Lys Glu Met Leu Pro Ile
 20 25 30

Val Asp Lys Pro Leu Ile Gln Tyr Ala Val Glu Glu Ala Val Glu Ala
 35 40 45

Gly Cys Thr Glu Met Val Phe Val Thr Gly Arg Asn Lys Arg Ser Ile
 50 55 60

Glu Asp His Phe Asp Lys Ala Tyr Glu Leu Glu Thr Glu Leu Glu Met
 65 70 75 80

Arg His Lys Asp Lys Leu Leu Glu His Val Arg Asn Ile Leu Pro Pro
 85 90 95

Asn Ile Thr Cys Leu Tyr Ile Arg Gln Ala Glu Ala Leu Gly Leu Gly
 100 105 110
 His Ala Val Leu Cys Ala Arg Ala Ala Ile Gly Asp Glu Pro Phe Ala
 115 120 125
 Val Ile Leu Ala Asp Asp Leu Ile Asp Ala Gln Lys Gly Ala Leu Lys
 130 135 140
 Gln Met Val Glu Val Tyr Glu Arg Ser Gly Asn Ser Ile Leu Gly Val
 145 150 155 160
 Glu Thr Val Glu Pro Ser Gln Thr Gly Ser Tyr Gly Ile Val Glu Thr
 165 170 175
 Glu Gln Leu Lys Gln Phe Gln Arg Ile Thr Gly Ile Val Glu Lys Pro
 180 185 190
 Lys Pro Glu Asp Ala Pro Ser Asn Leu Ala Val Val Gly Arg Tyr Ile
 195 200 205
 Leu Thr Pro Arg Ile Phe Asp Leu Leu Thr Gly Leu Pro Arg Gly Ala
 210 215 220
 Gly Asn Glu Ile Gln Leu Thr Asp Gly Ile Ala Lys Leu Leu Asp His
 225 230 235 240
 Glu Phe Val Leu Ala His Pro Phe Glu Gly Thr Arg Tyr Asp Cys Gly
 245 250 255
 Ser Lys Leu Gly Tyr Leu Glu Ala Thr Val Ala Tyr Gly Leu Lys His
 260 265 270
 Pro Glu Thr Gly Glu Pro Phe Arg Arg Leu Leu Glu Lys Tyr Arg Thr
 275 280 285
 Glu

<210> 155

<211> 876

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(876)

<400> 155

ttg acc gtc cga acg aaa aag acg gcg cat tat acc cta ttc cat tcc	48
Leu Thr Val Arg Thr Lys Lys Thr Ala His Tyr Thr Leu Phe His Ser	
1 5 10 15	
gac cga aaa ccg aac atg act act ctc aaa ccc gcc ctg ccc gct tat	96
Asp Arg Lys Pro Asn Met Thr Thr Leu Lys Pro Ala Leu Pro Ala Tyr	
20 25 30	
ctg gac aac atc cgc atc atc ctc acg cgc acc agc cat ccc gcc aac	144
Leu Asp Asn Ile Arg Ile Ile Leu Thr Arg Thr Ser His Pro Ala Asn	
35 40 45	
atc ggc tct gcc gcg cgc gcg atg aaa aca atg ggt ctg cac aaa ctg	192
Ile Gly Ser Ala Ala Arg Ala Met Lys Thr Met Gly Leu His Lys Leu	
50 55 60	
acc atc gtc gcc cca aat ctg atg gca acg ccg atg acg gaa aac ccg	240
Thr Ile Val Ala Pro Asn Leu Met Ala Thr Pro Met Thr Glu Asn Pro	
65 70 75 80	
ccc gtg ttt gac ccg gag cat cct caa tcg ttt aaa tta ccg gaa gaa	288
Pro Val Phe Asp Pro Glu His Pro Gln Ser Phe Lys Leu Pro Glu Glu	
85 90 95	
agc ttc atc ctc gct tcc ggc gcg gca gac gtt ttg gaa aat gcc acc	336
Ser Phe Ile Leu Ala Ser Gly Ala Ala Asp Val Leu Glu Asn Ala Thr	
100 105 110	
att gcc gct tct ttg gac gaa gcc ctt gcc gac acc acc atc gcc tgc	384
Ile Ala Ala Ser Leu Asp Glu Ala Leu Ala Asp Thr Thr Ile Ala Cys	
115 120 125	
gcc ctg acc agc cgc cgc cgc gaa att act gcg ccg ctg caa acc ccg	432
Ala Leu Thr Ser Arg Arg Arg Glu Ile Thr Ala Pro Leu Gln Thr Pro	
130 135 140	
cgc gat ttg gta tcc gaa tta ctg cag acc gca aac cga ggc gag aaa	480
Arg Asp Leu Val Ser Glu Leu Leu Gln Thr Ala Asn Arg Gly Glu Lys	
145 150 155 160	
gtg gca ctg gtt ttc ggc aac gag act ttc ggc ttg agc atc gaa gaa	528
Val Ala Leu Val Phe Gly Asn Glu Thr Phe Gly Leu Ser Ile Glu Glu	
165 170 175	
gtc caa gcc tgc aac cga ctg atg acc atc aac ggc aat ccc gac tat	576

Val Gln Ala Cys Asn Arg Leu Met Thr Ile Asn Gly Asn Pro Asp Tyr
 180 185 190
 ttc tcg ctc aac ctc gcc caa gcc gtg cag gtc gtg tgc tac gaa atc 624
 Phe Ser Leu Asn Leu Ala Gln Ala Val Gln Val Val Cys Tyr Glu Ile
 195 200 205
 ttc agc caa acc ggt tcg ccc atg acc cat ctt caa caa gaa gac cac 672
 Phe Ser Gln Thr Gly Ser Pro Met Thr His Leu Gln Gln Glu Asp His
 210 215 220
 gct gcg acc cac gag caa atc aaa ggc atg gtc gcc cac atg gaa agc 720
 Ala Ala Thr His Glu Gln Ile Lys Gly Met Val Ala His Met Glu Ser
 225 230 235 240
 gtg atg aac gac atc ggc ttt ttc aac cgc cgc aac ggc gag cgt ctg 768
 Val Met Asn Asp Ile Gly Phe Phe Asn Arg Arg Asn Gly Glu Arg Leu
 245 250 255
 atg cgc cgt atg cag agc ctg ttc gga cgc gcc aac acg caa acc gaa 816
 Met Arg Arg Met Gln Ser Leu Phe Gly Arg Ala Asn Thr Gln Thr Glu
 260 265 270
 gac atc gat atc ctg cgc ggt ttt ttc aat acc gtc agc cat cgt atc 864
 Asp Ile Asp Ile Leu Arg Gly Phe Phe Asn Thr Val Ser His Arg Ile
 275 280 285
 cat aaa aaa gac 876
 His Lys Lys Asp
 290
 <210> 156
 <211> 292
 <212> PRT
 <213> Neisseria meningitidis
 <400> 156
 Leu Thr Val Arg Thr Lys Lys Thr Ala His Tyr Thr Leu Phe His Ser
 1 5 10 15
 Asp Arg Lys Pro Asn Met Thr Thr Leu Lys Pro Ala Leu Pro Ala Tyr
 20 25 30
 Leu Asp Asn Ile Arg Ile Ile Leu Thr Arg Thr Ser His Pro Ala Asn
 35 40 45
 Ile Gly Ser Ala Ala Arg Ala Met Lys Thr Met Gly Leu His Lys Leu

50 55 60
 Thr Ile Val Ala Pro Asn Leu Met Ala Thr Pro Met Thr Glu Asn Pro
 65 70 75 80
 Pro Val Phe Asp Pro Glu His Pro Gln Ser Phe Lys Leu Pro Glu Glu
 85 90 95
 Ser Phe Ile Leu Ala Ser Gly Ala Ala Asp Val Leu Glu Asn Ala Thr
 100 105 110
 Ile Ala Ala Ser Leu Asp Glu Ala Leu Ala Asp Thr Thr Ile Ala Cys
 115 120 125
 Ala Leu Thr Ser Arg Arg Arg Glu Ile Thr Ala Pro Leu Gln Thr Pro
 130 135 140
 Arg Asp Leu Val Ser Glu Leu Leu Gln Thr Ala Asn Arg Gly Glu Lys
 145 150 155 160
 Val Ala Leu Val Phe Gly Asn Glu Thr Phe Gly Leu Ser Ile Glu Glu
 165 170 175
 Val Gln Ala Cys Asn Arg Leu Met Thr Ile Asn Gly Asn Pro Asp Tyr
 180 185 190
 Phe Ser Leu Asn Leu Ala Gln Ala Val Gln Val Val Cys Tyr Glu Ile
 195 200 205
 Phe Ser Gln Thr Gly Ser Pro Met Thr His Leu Gln Gln Glu Asp His
 210 215 220
 Ala Ala Thr His Glu Gln Ile Lys Gly Met Val Ala His Met Glu Ser
 225 230 235 240
 Val Met Asn Asp Ile Gly Phe Phe Asn Arg Arg Asn Gly Glu Arg Leu
 245 250 255
 Met Arg Arg Met Gln Ser Leu Phe Gly Arg Ala Asn Thr Gln Thr Glu
 260 265 270
 Asp Ile Asp Ile Leu Arg Gly Phe Phe Asn Thr Val Ser His Arg Ile
 275 280 285
 His Lys Lys Asp
 290

<210> 157

<211> 1008

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(1008)

<400> 157

atg tcc atc aaa atc ctg att att tcc ccc agt tgg ata ggc gac tgc 48
 Met Ser Ile Lys Ile Leu Ile Ile Ser Pro Ser Trp Ile Gly Asp Cys
 1 5 10 15

gtg atg acc cag ccc ttg ttc cgc cgt ttg aag gaa ctt cac ccc ggt 96
 Val Met Thr Gln Pro Leu Phe Arg Arg Leu Lys Glu Leu His Pro Gly
 20 25 30

tgc acg att gat gtg ttc gca ccg aag tgg tcg atg gcg gtg ttt gag 144
 Cys Thr Ile Asp Val Phe Ala Pro Lys Trp Ser Met Ala Val Phe Glu
 35 40 45

cgt atg ccg gaa gtg aat gag att ctt gaa aat ccg ttc gga cac ggt 192
 Arg Met Pro Glu Val Asn Glu Ile Leu Glu Asn Pro Phe Gly His Gly
 50 55 60

gcg ttg gag ttg aaa cgc cgt tgg cgg gtc ggt agg gat ttg ggg cgg 240
 Ala Leu Glu Leu Lys Arg Arg Trp Arg Val Gly Arg Asp Leu Gly Arg
 65 70 75 80

cgc gga tac gat cag gtt atc gtg ttg ccc ggt tct ttg aaa tcg gca 288
 Arg Gly Tyr Asp Gln Val Ile Val Leu Pro Gly Ser Leu Lys Ser Ala
 85 90 95

atc atc gcg ctg gcg aca ggt atc ggt aaa agg acg ggt tat gtc ggt 336
 Ile Ile Ala Leu Ala Thr Gly Ile Gly Lys Arg Thr Gly Tyr Val Gly
 100 105 110

gaa agc cgt tat ttt ctg ttg aac gat ata cgc agg ctg gat aag gaa 384
 Glu Ser Arg Tyr Phe Leu Leu Asn Asp Ile Arg Arg Leu Asp Lys Glu
 115 120 125

cgt ctg cct ttg atg gtg gac aga tat acg gct ctc gcg cat ccg agc 432
 Arg Leu Pro Leu Met Val Asp Arg Tyr Thr Ala Leu Ala His Pro Ser
 130 135 140

cag gag gat ttt gac ggg cat tcg gga ttc ccc gag ttt tcc att gat 480

311

<210> 158

<211> 336

<212> PRT

<213> Neisseria meningitidis

<400> 158

Met Ser Ile Lys Ile Leu Ile Ile Ser Pro Ser Trp Ile Gly Asp Cys
 1 5 10 15

Val Met Thr Gln Pro Leu Phe Arg Arg Leu Lys Glu Leu His Pro Gly
 20 25 30

Cys Thr Ile Asp Val Phe Ala Pro Lys Trp Ser Met Ala Val Phe Glu
 35 40 45

Arg Met Pro Glu Val Asn Glu Ile Leu Glu Asn Pro Phe Gly His Gly
 50 55 60

Ala Leu Glu Leu Lys Arg Arg Trp Arg Val Gly Arg Asp Leu Gly Arg
 65 70 75 80

Arg Gly Tyr Asp Gln Val Ile Val Leu Pro Gly Ser Leu Lys Ser Ala
 85 90 95

Ile Ile Ala Leu Ala Thr Gly Ile Gly Lys Arg Thr Gly Tyr Val Gly
 100 105 110

Glu Ser Arg Tyr Phe Leu Leu Asn Asp Ile Arg Arg Leu Asp Lys Glu
 115 120 125

Arg Leu Pro Leu Met Val Asp Arg Tyr Thr Ala Leu Ala His Pro Ser
 130 135 140

Gln Glu Asp Phe Asp Gly His Ser Gly Phe Pro Glu Phe Ser Ile Asp
 145 150 155 160

Glu Arg Arg Arg Glu Ile Ser Val Glu Thr Phe Gly Leu Asp Ile Gly
 165 170 175

Lys Pro Val Leu Ala Phe Cys Pro Gly Ala Glu Phe Gly Pro Ala Lys
 180 185 190

Arg Trp Pro Thr Arg Tyr Phe Ala Glu Leu Gly Lys His Tyr Ser Ala
 195 200 205

Ala Gly Trp Gln Val Trp Leu Phe Gly Ser Gln Lys Asp Asp Glu Ile
 210 215 220

Ala Glu Glu Ile Asn Cys Leu Ser Asp Gly Met Cys Val Asn Leu Cys
 225 230 235 240

Gly Lys Thr Asp Leu Ser Gln Ala Met Asp Leu Leu Ser Leu Ala Asp
 245 250 255

Thr Val Val Cys Asn Asp Ser Gly Leu Met His Leu Ala Ala Ala Leu
 260 265 270

Gly Arg Lys Val Val Ala Val Tyr Gly Ser Ser Ser Pro Thr His Thr
 275 280 285

Pro Pro Leu Ser Asp Arg Ala Lys Ile Val Ser Leu His Leu Glu Cys
 290 295 300

Ser Pro Cys Phe Lys Arg Glu Cys Pro Leu Gly His Thr Asp Cys Leu
 305 310 315 320

Asn Arg Leu Tyr Pro Glu Lys Ile Val Gln Ala Val Glu Glu Ala Val
 325 330 335

<210> 159

<211> 381

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(381)

<400> 159

atg gaa cct tcc tcc tac gcg gca gaa aaa aaa gga aaa ggc ggc atc 48
 Met Glu Pro Ser Ser Tyr Ala Ala Glu Lys Lys Gly Lys Gly Gly Ile
 1 5 10 15

agg cgc gtc atc aac gca ttc ggc tat tcg ata gac ggc atc gcc gcc 96
 Arg Arg Val Ile Asn Ala Phe Gly Tyr Ser Ile Asp Gly Ile Ala Ala
 20 25 30

gcc tac cgt tac gaa gcg gca ttc cgt cag gtt ttg tgg ctg aac gcg 144
 Ala Tyr Arg Tyr Glu Ala Ala Phe Arg Gln Val Leu Trp Leu Asn Ala
 35 40 45

ctg ctg gtg tgc gcg gca ttt ttt tgg gtt tcc gaa aag tcc ctc cgc 192
 Leu Leu Val Cys Ala Ala Phe Phe Trp Val Ser Glu Lys Ser Leu Arg
 50 55 60

ctg ccg ttg att atc gcg tct ttt gtg tgc gtc att gtc gaa ctg ttc 240
 Leu Pro Leu Ile Ile Ala Ser Phe Val Ser Val Ile Val Glu Leu Phe
 65 70 75 80

aac act gcc gtc gaa gcc gcc gtc gat cat act tcg act gaa aaa cac 288
 Asn Thr Ala Val Glu Ala Ala Val Asp His Thr Ser Thr Glu Lys His
 85 90 95

gag ctg gct aaa cgc gcc aaa gac gca ggt tct gct gca caa ttg ttc 336
 Glu Leu Ala Lys Arg Ala Lys Asp Ala Gly Ser Ala Ala Gln Leu Phe
 100 105 110

gcg atg ctg atg tta gcg gcg gtt tgg ctg tcc gcc ctg ttc ggg 381
 Ala Met Leu Met Leu Ala Ala Val Trp Leu Ser Ala Leu Phe Gly
 115 120 125

<210> 160

<211> 127

<212> PRT

<213> Neisseria meningitidis

<400> 160

Met Glu Pro Ser Ser Tyr Ala Ala Glu Lys Lys Gly Lys Gly Gly Ile
 1 5 10 15

Arg Arg Val Ile Asn Ala Phe Gly Tyr Ser Ile Asp Gly Ile Ala Ala
 20 25 30

Ala Tyr Arg Tyr Glu Ala Ala Phe Arg Gln Val Leu Trp Leu Asn Ala
 35 40 45

Leu Leu Val Cys Ala Ala Phe Phe Trp Val Ser Glu Lys Ser Leu Arg
 50 55 60

Leu Pro Leu Ile Ile Ala Ser Phe Val Ser Val Ile Val Glu Leu Phe
 65 70 75 80

Asn Thr Ala Val Glu Ala Ala Val Asp His Thr Ser Thr Glu Lys His
 85 90 95

Glu Leu Ala Lys Arg Ala Lys Asp Ala Gly Ser Ala Ala Gln Leu Phe
 100 105 110

Ala Met Leu Met Leu Ala Ala Val Trp Leu Ser Ala Leu Phe Gly
 115 120 125

<210> 161
 <211> 990
 <212> DNA
 <213> *Neisseria meningitidis*

<220>
 <221> CDS
 <222> (1)..(990)

<400> 161
 atg aaa aaa gaa agc cgc ccc gcg ttt gac gcg gca gcc gta ttt gac 48
 Met Lys Lys Glu Ser Arg Pro Ala Phe Asp Ala Ala Ala Val Phe Asp
 1 5 10 15
 gcg gca gcc gta ccg gta tcc gac agc ggg ttt gcc gcc aat gca aat 96
 Ala Ala Ala Val Pro Val Ser Asp Ser Gly Phe Ala Ala Asn Ala Asn
 20 25 30
 gtc cgc cgt ttt gtg gac gat gaa gtc ggg aaa ggg gat ttt tcc cgg 144
 Val Arg Arg Phe Val Asp Asp Glu Val Gly Lys Gly Asp Phe Ser Arg
 35 40 45
 gcg gaa tgg cag gat ttt ttt gac aaa gcg gct tac aag gcg gac atc 192
 Ala Glu Trp Gln Asp Phe Phe Asp Lys Ala Ala Tyr Lys Ala Asp Ile
 50 55 60
 gtc aag att atg cac cgc ccc tcc aca tcg cgt ccg tgg tat gtg ttc 240
 Val Lys Ile Met His Arg Pro Ser Thr Ser Arg Pro Trp Tyr Val Phe
 65 70 75 80
 cgc acg gga aat tcg ggc aag gcg aaa ttt cgc ggc gcg cgc cgg ttt 288
 Arg Thr Gly Asn Ser Gly Lys Ala Lys Phe Arg Gly Ala Arg Arg Phe
 85 90 95
 tat gcg gaa aac cgc gcg ctt atc gat gat gtg gcg caa aaa tac ggc 336
 Tyr Ala Glu Asn Arg Ala Leu Ile Asp Asp Val Ala Gln Lys Tyr Gly
 100 105 110
 gtg cct gcc gaa ctt atc gtg gcg gtt atc ggg att gaa acg aat tac 384
 Val Pro Ala Glu Leu Ile Val Ala Val Ile Gly Ile Glu Thr Asn Tyr
 115 120 125
 ggc aaa aat acg ggc agt ttc cgt gtg gcg gac gca ttg gcg acc tta 432
 Gly Lys Asn Thr Gly Ser Phe Arg Val Ala Asp Ala Leu Ala Thr Leu
 130 135 140

ggc ttt gat tac ccc cgc cgc gcc ggg ttt ttc caa aaa gaa ttg gtc 480
 Gly Phe Asp Tyr Pro Arg Arg Ala Gly Phe Phe Gln Lys Glu Leu Val
 145 150 155 160

gag ctt tta aag ctg gca aaa gaa gaa ggc ggc gat gtt ttc gcc ttt 528
 Glu Leu Leu Lys Leu Ala Lys Glu Glu Gly Gly Asp Val Phe Ala Phe
 165 170 175

aaa ggc agc tat gcg ggc gca atg ggg atg ccg caa ttt atg cct tcg 576
 Lys Gly Ser Tyr Ala Gly Ala Met Gly Met Pro Gln Phe Met Pro Ser
 180 185 190

agc tac cgg aaa tgg gcg gtg gat tat gac ggg gac gga cat cgg gac 624
 Ser Tyr Arg Lys Trp Ala Val Asp Tyr Asp Gly Asp Gly His Arg Asp
 195 200 205

ata tgg ggc aat gtt ggc gat gtc gcg gca tcg att gcc aat tat atg 672
 Ile Trp Gly Asn Val Gly Asp Val Ala Ala Ser Ile Ala Asn Tyr Met
 210 215 220

aag cag cac ggt tgg cgc acg ggc ggg aaa ata ctg gtg tct gca aca 720
 Lys Gln His Gly Trp Arg Thr Gly Gly Lys Ile Leu Val Ser Ala Thr
 225 230 235 240

ttg gcg ccg ggt gcg gat gtt cag gca atc att ggc gaa aaa acc gcc 768
 Leu Ala Pro Gly Ala Asp Val Gln Ala Ile Ile Gly Glu Lys Thr Ala
 245 250 255

ctg acg cgg acg gtg gcg gat ttg aag gcg tac ggc atc atc ccc ggc 816
 Leu Thr Arg Thr Val Ala Asp Leu Lys Ala Tyr Gly Ile Ile Pro Gly
 260 265 270

gaa gag ctt gcc gat gat gaa aag gcg gtt ttg ttc aaa ctg gaa acc 864
 Glu Glu Leu Ala Asp Asp Glu Lys Ala Val Leu Phe Lys Leu Glu Thr
 275 280 285

gca ccc ggc gtg ttt gaa tat tat ttg ggc ttg aac aat ttt tat acg 912
 Ala Pro Gly Val Phe Glu Tyr Tyr Leu Gly Leu Asn Asn Phe Tyr Thr
 290 295 300

gta tgg cag tac aat cac agt cgg atg tat gta acg gcg gtc agg gac 960
 Val Trp Gln Tyr Asn His Ser Arg Met Tyr Val Thr Ala Val Arg Asp
 305 310 315 320

att gcc aat tcg ctt ggc ggc ccg gga ttg 990
 Ile Ala Asn Ser Leu Gly Gly Pro Gly Leu
 325 330

<210> 162

<211> 330

<212> PRT

<213> *Neisseria meningitidis*

<400> 162

Met Lys Lys Glu Ser Arg Pro Ala Phe Asp Ala Ala Ala Val Phe Asp
 1 5 10 15

Ala Ala Ala Val Pro Val Ser Asp Ser Gly Phe Ala Ala Asn Ala Asn
 20 25 30

Val Arg Arg Phe Val Asp Asp Glu Val Gly Lys Gly Asp Phe Ser Arg
 35 40 45

Ala Glu Trp Gln Asp Phe Phe Asp Lys Ala Ala Tyr Lys Ala Asp Ile
 50 55 60

Val Lys Ile Met His Arg Pro Ser Thr Ser Arg Pro Trp Tyr Val Phe
 65 70 75 80

Arg Thr Gly Asn Ser Gly Lys Ala Lys Phe Arg Gly Ala Arg Arg Phe
 85 90 95

Tyr Ala Glu Asn Arg Ala Leu Ile Asp Asp Val Ala Gln Lys Tyr Gly
 100 105 110

Val Pro Ala Glu Leu Ile Val Ala Val Ile Gly Ile Glu Thr Asn Tyr
 115 120 125

Gly Lys Asn Thr Gly Ser Phe Arg Val Ala Asp Ala Leu Ala Thr Leu
 130 135 140

Gly Phe Asp Tyr Pro Arg Arg Ala Gly Phe Phe Gln Lys Glu Leu Val
 145 150 155 160

Glu Leu Leu Lys Leu Ala Lys Glu Glu Gly Gly Asp Val Phe Ala Phe
 165 170 175

Lys Gly Ser Tyr Ala Gly Ala Met Gly Met Pro Gln Phe Met Pro Ser
 180 185 190

Ser Tyr Arg Lys Trp Ala Val Asp Tyr Asp Gly Asp Gly His Arg Asp
 195 200 205

Ile Trp Gly Asn Val Gly Asp Val Ala Ala Ser Ile Ala Asn Tyr Met
 210 215 220

Lys Gln His Gly Trp Arg Thr Gly Gly Lys Ile Leu Val Ser Ala Thr
 225 230 235 240

Leu Ala Pro Gly Ala Asp Val Gln Ala Ile Ile Gly Glu Lys Thr Ala
 245 250 255

Leu Thr Arg Thr Val Ala Asp Leu Lys Ala Tyr Gly Ile Ile Pro Gly
 260 265 270

Glu Glu Leu Ala Asp Asp Glu Lys Ala Val Leu Phe Lys Leu Glu Thr
 275 280 285

Ala Pro Gly Val Phe Glu Tyr Tyr Leu Gly Leu Asn Asn Phe Tyr Thr
 290 295 300

Val Trp Gln Tyr Asn His Ser Arg Met Tyr Val Thr Ala Val Arg Asp
 305 310 315 320

Ile Ala Asn Ser Leu Gly Gly Pro Gly Leu
 325 330

<210> 163

<211> 1773

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1773)

<400> 163

atg agt atc gtg ctg cac ggc gtg gcg gcg ggc aaa ggc att gcc gtc 48
 Met Ser Ile Val Leu His Gly Val Ala Ala Gly Lys Gly Ile Ala Val
 1 5 10 15

ggt tgc gcc cac ctg att gcg cgc ggt acg gag gaa gtg ccg cag tat 96
 Gly Cys Ala His Leu Ile Ala Arg Gly Thr Glu Glu Val Pro Gln Tyr
 20 25 30

gat gtt gcg gag gcg gac acc gat gcc gaa gcc gaa cgt ttc gat gcc 144
 Asp Val Ala Glu Ala Asp Thr Asp Ala Glu Ala Glu Arg Phe Asp Ala
 35 40 45

gcc gtc aaa gcc acg cgc aaa gag ttg gaa cag ctc cgc agc gcg att 192
 Ala Val Lys Ala Thr Arg Lys Glu Leu Glu Gln Leu Arg Ser Ala Ile

50	55	60	
ccc gaa aac gcc ccg acc gag ttg ggc gcg ttc atc tcg ctg cac ctg			240
Pro Glu Asn Ala Pro Thr Glu Leu Gly Ala Phe Ile Ser Leu His Leu			
65	70	75	80
atg ctc ttg acc gat gtt acc ttg tcg cgc gaa ccc gtc gat att tta			288
Met Leu Leu Thr Asp Val Thr Leu Ser Arg Glu Pro Val Asp Ile Leu			
	85	90	95
agg gaa caa aaa atc aac gcc gag tgg gca ttg aag cag cag agc gac			336
Arg Glu Gln Lys Ile Asn Ala Glu Trp Ala Leu Lys Gln Gln Ser Asp			
	100	105	110
aaa ctc gcc gcc caa ttc gac aat atg gac gat gcc tat ttg cgc gaa			384
Lys Leu Ala Ala Gln Phe Asp Asn Met Asp Asp Ala Tyr Leu Arg Glu			
	115	120	125
cgc aag cag gat atg ctg caa gtc gtc cgc cgc atc cac aac aac ctg			432
Arg Lys Gln Asp Met Leu Gln Val Val Arg Arg Ile His Asn Asn Leu			
	130	135	140
atc ggg cag ggc aac gag ttg gaa gtt gcc gac aac ctg ttt gac gaa			480
Ile Gly Gln Gly Asn Glu Leu Glu Val Ala Asp Asn Leu Phe Asp Glu			
	145	150	155
acc gtt ctg att gca aac gac ctt tcg ccc gcc gac acg gtt ttg ttt			528
Thr Val Leu Ile Ala Asn Asp Leu Ser Pro Ala Asp Thr Val Leu Phe			
	165	170	175
aaa gag cag cgc att gcc gcc ttc gtt acc gat gcc ggc ggc ccc acc			576
Lys Glu Gln Arg Ile Ala Ala Phe Val Thr Asp Ala Gly Gly Pro Thr			
	180	185	190
ggg cat acg gcg att ttg ggc agg agc ttg gac atc ccg tcc gtc gtc			624
Gly His Thr Ala Ile Leu Gly Arg Ser Leu Asp Ile Pro Ser Val Val			
	195	200	205
ggg ctg cac aac gcg cgc aaa ctg att acc gag ggc gaa acg gtc att			672
Gly Leu His Asn Ala Arg Lys Leu Ile Thr Glu Gly Glu Thr Val Ile			
	210	215	220
gtg gac ggt atc aac ggc gtg ttg att atc gcg ccg gat gag tcg gtg			720
Val Asp Gly Ile Asn Gly Val Leu Ile Ile Ala Pro Asp Glu Ser Val			
	225	230	235
ttg aac gaa tac cgc cgc cgt gcc cgc gaa tac cgc agc cac aaa cgc			768
Leu Asn Glu Tyr Arg Arg Arg Ala Arg Glu Tyr Arg Ser His Lys Arg			

245	250	255	
gat ttg aac aag ctc aaa aaa acc gcc gcc gcc acc gct gac ggg gtc			816
Asp Leu Asn Lys Leu Lys Lys Thr Ala Ala Ala Thr Ala Asp Gly Val			
260	265	270	
tgç atc gag ctt gtg ggc aat ata gaa tcc gcc gaa gac gtg aaa ccg			864
Cys Ile Glu Leu Val Gly Asn Ile Glu Ser Ala Glu Asp Val Lys Pro			
275	280	285	
ctg cac aac ctc ggc gca gac ggc atc ggg ctg ttc cgc agc gag ttt			912
Leu His Asn Leu Gly Ala Asp Gly Ile Gly Leu Phe Arg Ser Glu Phe			
290	295	300	
ctt tac ctg aac cgc gat acg atg ccg tct gaa gac gag cag tac gaa			960
Leu Tyr Leu Asn Arg Asp Thr Met Pro Ser Glu Asp Glu Gln Tyr Glu			
305	310	315	320
gtg tac agc gcg att gtc aaa aaa atg aaa ggc aaa agc gta acg ata			1008
Val Tyr Ser Ala Ile Val Lys Lys Met Lys Gly Lys Ser Val Thr Ile			
325	330	335	
cgg aca gtc gat ttg ggt gtg gac aaa aac ccg cgc tgg ttc ggg aaa			1056
Arg Thr Val Asp Leu Gly Val Asp Lys Asn Pro Arg Trp Phe Gly Lys			
340	345	350	
aac agc acg ccc aac ggc agc ctc aac ccc gcg ctg ggc atg acc ggc			1104
Asn Ser Thr Pro Asn Gly Ser Leu Asn Pro Ala Leu Gly Met Thr Gly			
355	360	365	
atc cgc ctg tgc ctt gcc gaa ccg gtc atg ttc cgc acc cag atg cgc			1152
Ile Arg Leu Cys Leu Ala Glu Pro Val Met Phe Arg Thr Gln Met Arg			
370	375	380	
gcc atc ctc cgc gcc gcc gcc cac ggc ccc gtg cgg atg atg tgg ccg			1200
Ala Ile Leu Arg Ala Ala Ala His Gly Pro Val Arg Met Met Trp Pro			
385	390	395	400
atg att acc tcc gta tcc gaa gtg cgc cag tgc ctc atc cac ctc gac			1248
Met Ile Thr Ser Val Ser Glu Val Arg Gln Cys Leu Ile His Leu Asp			
405	410	415	
acc gcg caa cgc cag ctt gcc gaa cgc ggc gat gcc ttc ggt aaa gtc			1296
Thr Ala Gln Arg Gln Leu Ala Glu Arg Gly Asp Ala Phe Gly Lys Val			
420	425	430	
ggc atc ggc tgt atg att gaa att ccg tct gcc gcg ctg acc gtc ggc			1344
Gly Ile Gly Cys Met Ile Glu Ile Pro Ser Ala Ala Leu Thr Val Gly			

435	440	445	
agt att ttg aaa ctg gtc gat ttc atc tcc gtc ggt acc aac gac ctg			1392
Ser Ile Leu Lys Leu Val Asp Phe Ile Ser Val Gly Thr Asn Asp Leu			
450	455	460	
att caa tac atc ttg tcc gtc gat cgc ggc gac gac agc gtc agc cac			1440
Ile Gln Tyr Ile Leu Ser Val Asp Arg Gly Asp Asp Ser Val Ser His			
465	470	475	480
ctc tac cag ccc ggc cat ccc gcc gtg ctg aaa atg ctg caa cac gtc			1488
Leu Tyr Gln Pro Gly His Pro Ala Val Leu Lys Met Leu Gln His Val			
	485	490	495
atc cgt acc gcc aac cgc atg gac aaa gac gta tcc gta tgc ggc gag			1536
Ile Arg Thr Ala Asn Arg Met Asp Lys Asp Val Ser Val Cys Gly Glu			
500	505	510	
atg gcg ggc gat acc gcg ttt acc cgc gtt tta ttg ggt atg ggg ctg			1584
Met Ala Gly Asp Thr Ala Phe Thr Arg Val Leu Leu Gly Met Gly Leu			
515	520	525	
cgc cgt ttt tcc atg aac ccc aac aac atc ctg ccc gtc aaa aac atc			1632
Arg Arg Phe Ser Met Asn Pro Asn Asn Ile Leu Pro Val Lys Asn Ile			
530	535	540	
att ctg cac agc aat gtc gga cag ctc gaa agc gat att gtg aaa gtc			1680
Ile Leu His Ser Asn Val Gly Gln Leu Glu Ser Asp Ile Val Lys Val			
545	550	555	560
atc cgc tgc gaa gac gaa gag aag tcg gaa aag ctg atc aaa cag atg			1728
Ile Arg Cys Glu Asp Glu Glu Lys Ser Glu Lys Leu Ile Lys Gln Met			
	565	570	575
aac agc gtg tct gtc gag gaa gaa gcc gac ttc aag ggg cgg aaa			1773
Asn Ser Val Ser Val Glu Glu Glu Ala Asp Phe Lys Gly Arg Lys			
580	585	590	

<210> 164

<211> 591

<212> PRT

<213> Neisseria meningitidis

<400> 164

Met Ser Ile Val Leu His Gly Val Ala Ala Gly Lys Gly Ile Ala Val
 1 5 10 15

Gly Cys Ala His Leu Ile Ala Arg Gly Thr Glu Glu Val Pro Gln Tyr
 20 25 30

Asp Val Ala Glu Ala Asp Thr Asp Ala Glu Ala Glu Arg Phe Asp Ala
 35 40 45

Ala Val Lys Ala Thr Arg Lys Glu Leu Glu Gln Leu Arg Ser Ala Ile
 50 55 60

Pro Glu Asn Ala Pro Thr Glu Leu Gly Ala Phe Ile Ser Leu His Leu
 65 70 75 80

Met Leu Leu Thr Asp Val Thr Leu Ser Arg Glu Pro Val Asp Ile Leu
 85 90 95

Arg Glu Gln Lys Ile Asn Ala Glu Trp Ala Leu Lys Gln Gln Ser Asp
 100 105 110

Lys Leu Ala Ala Gln Phe Asp Asn Met Asp Asp Ala Tyr Leu Arg Glu
 115 120 125

Arg Lys Gln Asp Met Leu Gln Val Val Arg Arg Ile His Asn Asn Leu
 130 135 140

Ile Gly Gln Gly Asn Glu Leu Glu Val Ala Asp Asn Leu Phe Asp Glu
 145 150 155 160

Thr Val Leu Ile Ala Asn Asp Leu Ser Pro Ala Asp Thr Val Leu Phe
 165 170 175

Lys Glu Gln Arg Ile Ala Ala Phe Val Thr Asp Ala Gly Gly Pro Thr
 180 185 190

Gly His Thr Ala Ile Leu Gly Arg Ser Leu Asp Ile Pro Ser Val Val
 195 200 205

Gly Leu His Asn Ala Arg Lys Leu Ile Thr Glu Gly Glu Thr Val Ile
 210 215 220

Val Asp Gly Ile Asn Gly Val Leu Ile Ile Ala Pro Asp Glu Ser Val
 225 230 235 240

Leu Asn Glu Tyr Arg Arg Arg Ala Arg Glu Tyr Arg Ser His Lys Arg
 245 250 255

Asp Leu Asn Lys Leu Lys Lys Thr Ala Ala Ala Thr Ala Asp Gly Val
 260 265 270

Cys Ile Glu Leu Val Gly Asn Ile Glu Ser Ala Glu Asp Val Lys Pro
 275 280 285

Leu His Asn Leu Gly Ala Asp Gly Ile Gly Leu Phe Arg Ser Glu Phe
 290 295 300

Leu Tyr Leu Asn Arg Asp Thr Met Pro Ser Glu Asp Glu Gln Tyr Glu
 305 310 315 320

Val Tyr Ser Ala Ile Val Lys Lys Met Lys Gly Lys Ser Val Thr Ile
 325 330 335

Arg Thr Val Asp Leu Gly Val Asp Lys Asn Pro Arg Trp Phe Gly Lys
 340 345 350

Asn Ser Thr Pro Asn Gly Ser Leu Asn Pro Ala Leu Gly Met Thr Gly
 355 360 365

Ile Arg Leu Cys Leu Ala Glu Pro Val Met Phe Arg Thr Gln Met Arg
 370 375 380

Ala Ile Leu Arg Ala Ala Ala His Gly Pro Val Arg Met Met Trp Pro
 385 390 395 400

Met Ile Thr Ser Val Ser Glu Val Arg Gln Cys Leu Ile His Leu Asp
 405 410 415

Thr Ala Gln Arg Gln Leu Ala Glu Arg Gly Asp Ala Phe Gly Lys Val
 420 425 430

Gly Ile Gly Cys Met Ile Glu Ile Pro Ser Ala Ala Leu Thr Val Gly
 435 440 445

Ser Ile Leu Lys Leu Val Asp Phe Ile Ser Val Gly Thr Asn Asp Leu
 450 455 460

Ile Gln Tyr Ile Leu Ser Val Asp Arg Gly Asp Asp Ser Val Ser His
 465 470 475 480

Leu Tyr Gln Pro Gly His Pro Ala Val Leu Lys Met Leu Gln His Val
 485 490 495

Ile Arg Thr Ala Asn Arg Met Asp Lys Asp Val Ser Val Cys Gly Glu
 500 505 510

Met Ala Gly Asp Thr Ala Phe Thr Arg Val Leu Leu Gly Met Gly Leu
 515 520 525

Arg Arg Phe Ser Met Asn Pro Asn Asn Ile Leu Pro Val Lys Asn Ile
530 535 540

Ile Leu His Ser Asn Val Gly Gln Leu Glu Ser Asp Ile Val Lys Val
545 550 555 560

Ile Arg Cys Glu Asp Glu Glu Lys Ser Glu Lys Leu Ile Lys Gln Met
565 570 575

Asn Ser Val Ser Val Glu Glu Glu Ala Asp Phe Lys Gly Arg Lys
580 585 590

<210> 165

<211> 381

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(381)

<400> 165

atg gaa cct tcc tcc tac gcg gca gaa aaa aaa gga aaa ggc ggc atc 48
Met Glu Pro Ser Ser Tyr Ala Ala Glu Lys Lys Gly Lys Gly Gly Ile
1 5 10 15

agg cgc gtc atc aac gca ttc ggc tat tcg ata gac ggc atc gcc gcc 96
Arg Arg Val Ile Asn Ala Phe Gly Tyr Ser Ile Asp Gly Ile Ala Ala
20 25 30

gcc tac cgt tac gaa gcg gca ttc cgt cag gtt ttg tgg ctg aac gcg 144
Ala Tyr Arg Tyr Glu Ala Ala Phe Arg Gln Val Leu Trp Leu Asn Ala
35 40 45

ctg ctg gtg tgc gcg gca ttt ttt tgg gtt tcc gaa aag tcc ctc cgc 192
Leu Leu Val Cys Ala Ala Phe Phe Trp Val Ser Glu Lys Ser Leu Arg
50 55 60

ctg ccg ttg att atc gcg tct ttt gtg tcg gtc att gtc gaa ctg ttc 240
Leu Pro Leu Ile Ile Ala Ser Phe Val Ser Val Ile Val Glu Leu Phe
65 70 75 80

aac act gcc gtc gaa gcc gcc gtc gat cat act tcg act gaa aaa cac 288
Asn Thr Ala Val Glu Ala Ala Val Asp His Thr Ser Thr Glu Lys His
85 90 95

gag ctg gct aaa cgc gcc aaa gac gca ggt tct gct gca caa ttg ttc 336
 Glu Leu Ala Lys Arg Ala Lys Asp Ala Gly Ser Ala Ala Gln Leu Phe
 100 105 110

gcg atg ctg atg tta gcg gcg gtt tgg ctg tcc gcc ctg ttc ggg 381
 Ala Met Leu Met Leu Ala Ala Val Trp Leu Ser Ala Leu Phe Gly
 115 120 125

<210> 166

<211> 127

<212> PRT

<213> Neisseria meningitidis

<400> 166

Met Glu Pro Ser Ser Tyr Ala Ala Glu Lys Lys Gly Lys Gly Gly Ile
 1 5 10 15

Arg Arg Val Ile Asn Ala Phe Gly Tyr Ser Ile Asp Gly Ile Ala Ala
 20 25 30

Ala Tyr Arg Tyr Glu Ala Ala Phe Arg Gln Val Leu Trp Leu Asn Ala
 35 40 45

Leu Leu Val Cys Ala Ala Phe Phe Trp Val Ser Glu Lys Ser Leu Arg
 50 55 60

Leu Pro Leu Ile Ile Ala Ser Phe Val Ser Val Ile Val Glu Leu Phe
 65 70 75 80

Asn Thr Ala Val Glu Ala Ala Val Asp His Thr Ser Thr Glu Lys His
 85 90 95

Glu Leu Ala Lys Arg Ala Lys Asp Ala Gly Ser Ala Ala Gln Leu Phe
 100 105 110

Ala Met Leu Met Leu Ala Ala Val Trp Leu Ser Ala Leu Phe Gly
 115 120 125

<210> 167

<211> 1257

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1257)

<400> 167

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atg agt atg gca ctt gcc caa aaa ctt gcc gcc gac agc att gcg gcg 48
Met Ser Met Ala Leu Ala Gln Lys Leu Ala Ala Asp Ser Ile Ala Ala
  1             5             10            15

gtt gcc gaa gga cgc aac ctt cag gac gtg ttg gcg caa atc cgc acc 96
Val Ala Glu Gly Arg Asn Leu Gln Asp Val Leu Ala Gln Ile Arg Thr
      20             25            30

gcg cat ccc gac ctt acg gcg cag gaa aac ggc gcg ttg cag gac atc 144
Ala His Pro Asp Leu Thr Ala Gln Glu Asn Gly Ala Leu Gln Asp Ile
      35             40            45

gcc tac ggc tgc cag cgt tat ttg ggc agt ttg aaa cat atg ctc gcg 192
Ala Tyr Gly Cys Gln Arg Tyr Leu Gly Ser Leu Lys His Met Leu Ala
      50             55            60

cag atg ctg aaa aag ccg att ggc aat ccg cag ctc gaa agt ctg ctt 240
Gln Met Leu Lys Lys Pro Ile Gly Asn Pro Gln Leu Glu Ser Leu Leu
      65             70            75            80

ttg gcg gcg ttg tac caa ctg cat tac acg cgc aac gcg ccg cat gct 288
Leu Ala Ala Leu Tyr Gln Leu His Tyr Thr Arg Asn Ala Pro His Ala
      85             90            95

gtg gtc aac gaa gct gtt gaa agc atc gcc aaa atc gga cgc ggg cag 336
Val Val Asn Glu Ala Val Glu Ser Ile Ala Lys Ile Gly Arg Gly Gln
      100            105            110

tac cgt tgc ttt gcc aac gcg att ttg cgc cgc ttt ttg cgc gaa cgc 384
Tyr Arg Ser Phe Ala Asn Ala Ile Leu Arg Arg Phe Leu Arg Glu Arg
      115            120            125

gac aag ctt gcg gct tcc tgc aaa aaa gac gat gtg gcg aaa cac aac 432
Asp Lys Leu Ala Ala Ser Cys Lys Lys Asp Asp Val Ala Lys His Asn
      130            135            140

ctg ccg ctg tgg tgg gtg gct tac ttg aaa aac cat tat ccg aaa cac 480
Leu Pro Leu Trp Trp Val Ala Tyr Leu Lys Asn His Tyr Pro Lys His
      145            150            155            160

tgg cac aac atc gcc gcc gcg ctg caa tcc cat ccg ccg atg acg ttg 528
Trp His Asn Ile Ala Ala Ala Leu Gln Ser His Pro Pro Met Thr Leu
      165            170            175

cgc gtc aac cgc cga cac ggc aac gcc gaa agc tat ttg gaa aaa ctg 576

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Arg Val Asn Arg Arg His Gly Asn Ala Glu Ser Tyr Leu Glu Lys Leu	
180 185 190	
gct gcg gaa ggc att gcg gca aag gcg ttg gac gaa tat gcg gtt acg	624
Ala Ala Glu Gly Ile Ala Ala Lys Ala Leu Asp Glu Tyr Ala Val Thr	
195 200 205	
ttg gaa gaa gcc gtg ccg gtg aac cgc ctg ccc ggt ttt tca gac ggc	672
Leu Glu Glu Ala Val Pro Val Asn Arg Leu Pro Gly Phe Ser Asp Gly	
210 215 220	
att gtt tcg gta cag gac ttc ggc gcg cag cag gcg gcg tat ctc ctt	720
Ile Val Ser Val Gln Asp Phe Gly Ala Gln Gln Ala Ala Tyr Leu Leu	
225 230 235 240	
aac ccg aaa gac ggc gaa cgg att ttg gac gcg tgc gcc gcg ccg ggc	768
Asn Pro Lys Asp Gly Glu Arg Ile Leu Asp Ala Cys Ala Ala Pro Gly	
245 250 255	
ggc aag acg ggg cat atc ttg gaa ctg gcg gat tgc cgc gtt acc gcc	816
Gly Lys Thr Gly His Ile Leu Glu Leu Ala Asp Cys Arg Val Thr Ala	
260 265 270	
ttg gac att gat gca ggc cgt ctg aag cgt gtg gag gac aat atc gcc	864
Leu Asp Ile Asp Ala Gly Arg Leu Lys Arg Val Glu Asp Asn Ile Ala	
275 280 285	
cgt ttg ggc ttt cag acg gca tcg gcg gcg tgt gcc gat gcg cgg gat	912
Arg Leu Gly Phe Gln Thr Ala Ser Ala Ala Cys Ala Asp Ala Arg Asp	
290 295 300	
ttg gcg gcg tgg tat gat gga aag cct ttc gac aca atc ctt gcc gac	960
Leu Ala Ala Trp Tyr Asp Gly Lys Pro Phe Asp Thr Ile Leu Ala Asp	
305 310 315 320	
gtg cca tgc acc gct tcc ggc gtg gcg cgg cgc aat ccc gac gtg aaa	1008
Val Pro Cys Thr Ala Ser Gly Val Ala Arg Arg Asn Pro Asp Val Lys	
325 330 335	
tgg ctg cgc cgt ccg acc gac gcg ctc aaa acc gcc cgc cag cag gaa	1056
Trp Leu Arg Arg Pro Thr Asp Ala Leu Lys Thr Ala Arg Gln Gln Glu	
340 345 350	
gcc ctg ctg gac gca ctt tgg cag acg ctg aaa cag ggc ggc aga atg	1104
Ala Leu Leu Asp Ala Leu Trp Gln Thr Leu Lys Gln Gly Gly Arg Met	
355 360 365	
ctg ctt gcc acc tgt tcc gtg ttc gtc gaa gaa aac gac ggg caa ttg	1152

Leu Leu Ala Thr Cys Ser Val Phe Val Glu Glu Asn Asp Gly Gln Leu
 370 375 380

caa aaa ttc ctc aac cgc cat gcc gat gcc gaa ccg atc gaa tcg cgg 1200
 Gln Lys Phe Leu Asn Arg His Ala Asp Ala Glu Pro Ile Glu Ser Arg
 385 390 395 400

gtg ctt tta ccg aac aaa cac caa gat ggc ttt tat tac gcg ctt att 1248
 Val Leu Leu Pro Asn Lys His Gln Asp Gly Phe Tyr Tyr Ala Leu Ile
 405 410 415

caa aag cat 1257
 Gln Lys His

<210> 168
 <211> 419
 <212> PRT
 <213> Neisseria meningitidis

<400> 168
 Met Ser Met Ala Leu Ala Gln Lys Leu Ala Ala Asp Ser Ile Ala Ala
 1 5 10 15

Val Ala Glu Gly Arg Asn Leu Gln Asp Val Leu Ala Gln Ile Arg Thr
 20 25 30

Ala His Pro Asp Leu Thr Ala Gln Glu Asn Gly Ala Leu Gln Asp Ile
 35 40 45

Ala Tyr Gly Cys Gln Arg Tyr Leu Gly Ser Leu Lys His Met Leu Ala
 50 55 60

Gln Met Leu Lys Lys Pro Ile Gly Asn Pro Gln Leu Glu Ser Leu Leu
 65 70 75 80

Leu Ala Ala Leu Tyr Gln Leu His Tyr Thr Arg Asn Ala Pro His Ala
 85 90 95

Val Val Asn Glu Ala Val Glu Ser Ile Ala Lys Ile Gly Arg Gly Gln
 100 105 110

Tyr Arg Ser Phe Ala Asn Ala Ile Leu Arg Arg Phe Leu Arg Glu Arg
 115 120 125

Asp Lys Leu Ala Ala Ser Cys Lys Lys Asp Asp Val Ala Lys His Asn
 130 135 140

Leu Pro Leu Trp Trp Val Ala Tyr Leu Lys Asn His Tyr Pro Lys His
 145 150 155 160

Trp His Asn Ile Ala Ala Ala Leu Gln Ser His Pro Pro Met Thr Leu
 165 170 175

Arg Val Asn Arg Arg His Gly Asn Ala Glu Ser Tyr Leu Glu Lys Leu
 180 185 190

Ala Ala Glu Gly Ile Ala Ala Lys Ala Leu Asp Glu Tyr Ala Val Thr
 195 200 205

Leu Glu Glu Ala Val Pro Val Asn Arg Leu Pro Gly Phe Ser Asp Gly
 210 215 220

Ile Val Ser Val Gln Asp Phe Gly Ala Gln Gln Ala Ala Tyr Leu Leu
 225 230 235 240

Asn Pro Lys Asp Gly Glu Arg Ile Leu Asp Ala Cys Ala Ala Pro Gly
 245 250 255

Gly Lys Thr Gly His Ile Leu Glu Leu Ala Asp Cys Arg Val Thr Ala
 260 265 270

Leu Asp Ile Asp Ala Gly Arg Leu Lys Arg Val Glu Asp Asn Ile Ala
 275 280 285

Arg Leu Gly Phe Gln Thr Ala Ser Ala Ala Cys Ala Asp Ala Arg Asp
 290 295 300

Leu Ala Ala Trp Tyr Asp Gly Lys Pro Phe Asp Thr Ile Leu Ala Asp
 305 310 315 320

Val Pro Cys Thr Ala Ser Gly Val Ala Arg Arg Asn Pro Asp Val Lys
 325 330 335

Trp Leu Arg Arg Pro Thr Asp Ala Leu Lys Thr Ala Arg Gln Gln Glu
 340 345 350

Ala Leu Leu Asp Ala Leu Trp Gln Thr Leu Lys Gln Gly Gly Arg Met
 355 360 365

Leu Leu Ala Thr Cys Ser Val Phe Val Glu Glu Asn Asp Gly Gln Leu
 370 375 380

Gln Lys Phe Leu Asn Arg His Ala Asp Ala Glu Pro Ile Glu Ser Arg
 385 390 395 400

Val Leu Leu Pro Asn Lys His Gln Asp Gly Phe Tyr Tyr Ala Leu Ile
 405 410 415

Gln Lys His

<210> 169

<211> 804

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(804)

<400> 169

atg aat atg aaa aaa tgg att gcc gcc gcc ctt gcc tgt tcc gcg ctc 48
 Met Asn Met Lys Lys Trp Ile Ala Ala Ala Leu Ala Cys Ser Ala Leu
 1 5 10 15

gcg ctg tct gcc tgc ggc ggt cag ggt aaa gat gcc gcc gcg ccc gcc 96
 Ala Leu Ser Ala Cys Gly Gly Gln Gly Lys Asp Ala Ala Ala Pro Ala
 20 25 30

gca aat ccc gac aaa gtg tac cgc gtg gct tcc aac gcc gag ttt gcc 144
 Ala Asn Pro Asp Lys Val Tyr Arg Val Ala Ser Asn Ala Glu Phe Ala
 35 40 45

ccc ttt gaa tct tta gac tcg aaa ggc aat gtc gaa ggt ttc gat gtg 192
 Pro Phe Glu Ser Leu Asp Ser Lys Gly Asn Val Glu Gly Phe Asp Val
 50 55 60

gat ttg atg aac gcg atg gcg aag gcg ggc aat ttt aaa atc gaa ttc 240
 Asp Leu Met Asn Ala Met Ala Lys Ala Gly Asn Phe Lys Ile Glu Phe
 65 70 75 80

aaa cac cag ccg tgg gac agc ctt ttc ccc gcc ttg aac aac ggc gat 288
 Lys His Gln Pro Trp Asp Ser Leu Phe Pro Ala Leu Asn Asn Gly Asp
 85 90 95

gcg gac gtt gtg atg tcg ggc gta acc att acc gac gac cgc aaa cag 336
 Ala Asp Val Val Met Ser Gly Val Thr Ile Thr Asp Asp Arg Lys Gln
 100 105 110

tct atg gac ttc agc gac ccg tat ttt gaa atc acc caa gtc gtc ctc 384
 Ser Met Asp Phe Ser Asp Pro Tyr Phe Glu Ile Thr Gln Val Val Leu

115	120	125	
gtt ccg aaa ggc aaa aaa ata tct tct tcc gaa gat ttg aaa aac atg			432
Val Pro Lys Gly Lys Lys Ile Ser Ser Ser Glu Asp Leu Lys Asn Met			
130	135	140	
aac aaa gtc ggc gtg gta acc ggc tac acg ggc gat ttc tcc gta tcc			480
Asn Lys Val Gly Val Val Thr Gly Tyr Thr Gly Asp Phe Ser Val Ser			
145	150	155	160
aaa ctc ttg ggc aac gac aac ccg aaa atc gcg cgc ttt gaa aac gtt			528
Lys Leu Leu Gly Asn Asp Asn Pro Lys Ile Ala Arg Phe Glu Asn Val			
165	170	175	
ccc ctg att atc aaa gaa ctg gaa aac ggc ggc ttg gat tcc gtg gtc			576
Pro Leu Ile Ile Lys Glu Leu Glu Asn Gly Gly Leu Asp Ser Val Val			
180	185	190	
agc gac agc gca gtc atc gcc aat tat gtg aaa aac aat ccg acc aaa			624
Ser Asp Ser Ala Val Ile Ala Asn Tyr Val Lys Asn Asn Pro Thr Lys			
195	200	205	
ggg atg gac ttc gtt acc ctg ccc gac ttc acc acc gaa cac tac ggc			672
Gly Met Asp Phe Val Thr Leu Pro Asp Phe Thr Thr Glu His Tyr Gly			
210	215	220	
atc gcg gta cgc aaa ggc gac gaa gca acc gtc aaa atg ctg aac gat			720
Ile Ala Val Arg Lys Gly Asp Glu Ala Thr Val Lys Met Leu Asn Asp			
225	230	235	240
gcg ttg aaa aaa gta cgc gaa agc ggc gaa tac gac aaa atc tac gcc			768
Ala Leu Lys Lys Val Arg Glu Ser Gly Glu Tyr Asp Lys Ile Tyr Ala			
245	250	255	
aaa tat ttt gca aaa gaa gac gga cag gcc gca aaa			804
Lys Tyr Phe Ala Lys Glu Asp Gly Gln Ala Ala Lys			
260	265		

<210> 170

<211> 268

<212> PRT

<213> Neisseria meningitidis

<400> 170

Met Asn Met Lys Lys Trp Ile Ala Ala Ala Leu Ala Cys Ser Ala Leu
1 5 10 15

Ala Leu Ser Ala Cys Gly Gly Gln Gly Lys Asp Ala Ala Ala Pro Ala
 20 25 30

Ala Asn Pro Asp Lys Val Tyr Arg Val Ala Ser Asn Ala Glu Phe Ala
 35 40 45

Pro Phe Glu Ser Leu Asp Ser Lys Gly Asn Val Glu Gly Phe Asp Val
 50 55 60

Asp Leu Met Asn Ala Met Ala Lys Ala Gly Asn Phe Lys Ile Glu Phe
 65 70 75 80

Lys His Gln Pro Trp Asp Ser Leu Phe Pro Ala Leu Asn Asn Gly Asp
 85 90 95

Ala Asp Val Val Met Ser Gly Val Thr Ile Thr Asp Asp Arg Lys Gln
 100 105 110

Ser Met Asp Phe Ser Asp Pro Tyr Phe Glu Ile Thr Gln Val Val Leu
 115 120 125

Val Pro Lys Gly Lys Lys Ile Ser Ser Ser Glu Asp Leu Lys Asn Met
 130 135 140

Asn Lys Val Gly Val Val Thr Gly Tyr Thr Gly Asp Phe Ser Val Ser
 145 150 155 160

Lys Leu Leu Gly Asn Asp Asn Pro Lys Ile Ala Arg Phe Glu Asn Val
 165 170 175

Pro Leu Ile Ile Lys Glu Leu Glu Asn Gly Gly Leu Asp Ser Val Val
 180 185 190

Ser Asp Ser Ala Val Ile Ala Asn Tyr Val Lys Asn Asn Pro Thr Lys
 195 200 205

Gly Met Asp Phe Val Thr Leu Pro Asp Phe Thr Thr Glu His Tyr Gly
 210 215 220

Ile Ala Val Arg Lys Gly Asp Glu Ala Thr Val Lys Met Leu Asn Asp
 225 230 235 240

Ala Leu Lys Lys Val Arg Glu Ser Gly Glu Tyr Asp Lys Ile Tyr Ala
 245 250 255

Lys Tyr Phe Ala Lys Glu Asp Gly Gln Ala Ala Lys
 260 265

<210> 171
 <211> 864
 <212> DNA
 <213> Neisseria meningitidis

<220>
 <221> CDS
 <222> (1)..(864)

<400> 171
 atg aaa gca aaa atc ctg act tcc gtt gca ctg ctt gcc tgt tcc ggc 48
 Met Lys Ala Lys Ile Leu Thr Ser Val Ala Leu Leu Ala Cys Ser Gly
 1 5 10 15

 agc ctg ttt gcc caa acg ctg gca acc gtc aac ggt cag aaa atc gac 96
 Ser Leu Phe Ala Gln Thr Leu Ala Thr Val Asn Gly Gln Lys Ile Asp
 20 25 30

 agt tcc gtc att gat gcg cag gtt gcc gca ttc cgt gcg gaa aac agc 144
 Ser Ser Val Ile Asp Ala Gln Val Ala Ala Phe Arg Ala Glu Asn Ser
 35 40 45

 cgt gcc gaa gac acg ccg caa ctg cgc caa tcc ctg ctg gaa aac gaa 192
 Arg Ala Glu Asp Thr Pro Gln Leu Arg Gln Ser Leu Leu Glu Asn Glu
 50 55 60

 gtg gtc aac acc gtg gtc gca cag gaa gtg aaa cgc ctg aaa ctc gac 240
 Val Val Asn Thr Val Val Ala Gln Glu Val Lys Arg Leu Lys Leu Asp
 65 70 75 80

 cgg tcg gca gag ttt aaa aat gcg ctt gcc aaa ttg cgt gcc gaa gcg 288
 Arg Ser Ala Glu Phe Lys Asn Ala Leu Ala Lys Leu Arg Ala Glu Ala
 85 90 95

 aaa aag tcg ggc gac gac aag aaa ccg tcc ttc aaa acc gtt tgg cag 336
 Lys Lys Ser Gly Asp Asp Lys Lys Pro Ser Phe Lys Thr Val Trp Gln
 100 105 110

 gcg gta aaa tat ggc ttg aac ggc gag gca tac gcg ctg cat atc gcc 384
 Ala Val Lys Tyr Gly Leu Asn Gly Glu Ala Tyr Ala Leu His Ile Ala
 115 120 125

 aaa acc caa ccg gtt tcc gag cag gaa gta aaa gcc gca tat gac aat 432
 Lys Thr Gln Pro Val Ser Glu Gln Glu Val Lys Ala Ala Tyr Asp Asn
 130 135 140

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atc agc ggt ttt tac aaa ggt acg cag gaa gtc cag ttg ggc gaa atc   480
Ile Ser Gly Phe Tyr Lys Gly Thr Gln Glu Val Gln Leu Gly Glu Ile
145                      150                      155                      160

ctg acc gac aag gaa gaa aat gca aaa aaa gcg gtt gcc gac ttg aag   528
Leu Thr Asp Lys Glu Glu Asn Ala Lys Lys Ala Val Ala Asp Leu Lys
                      165                      170                      175

gcg aaa aaa ggt ttc gat gcc gtc ttg aaa caa tat tcc ctc aac gac   576
Ala Lys Lys Gly Phe Asp Ala Val Leu Lys Gln Tyr Ser Leu Asn Asp
                      180                      185                      190

cgt acc aaa cag acc ggt gcg ccg gtc gga tat gtg ccg ctg aaa gat   624
Arg Thr Lys Lys Gln Thr Gly Ala Pro Val Gly Tyr Val Pro Leu Lys Asp
                      195                      200                      205

ttg gaa cag ggt gtt ccg ccg ctt tat cag gca att aag gac ttg aaa   672
Leu Glu Gln Gly Val Pro Pro Leu Tyr Gln Ala Ile Lys Asp Leu Lys
                      210                      215                      220

aaa ggc gaa ttt acg gca acg ccg ctg aaa aac ggc gat ttc tac ggc   720
Lys Gly Glu Phe Thr Ala Thr Pro Leu Lys Asn Gly Asp Phe Tyr Gly
225                      230                      235                      240

gtt tat tat gtc aac gac agc cgc gag gta aaa gtg cct tct ttt gat   768
Val Tyr Tyr Val Asn Asp Ser Arg Glu Val Lys Val Pro Ser Phe Asp
                      245                      250                      255

gaa atg aaa gga cag att gcg ggc aac ctt cag gcg gaa cgg att gac   816
Glu Met Lys Gly Gln Ile Ala Gly Asn Leu Gln Ala Glu Arg Ile Asp
                      260                      265                      270

cgt gcc gtc ggt gca ctg ttg ggc aag gca aac atc aaa cct gca aaa   864
Arg Ala Val Gly Ala Leu Leu Gly Lys Ala Asn Ile Lys Pro Ala Lys
                      275                      280                      285

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<210> 172

<211> 288

<212> PRT

<213> Neisseria meningitidis

<400> 172

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Met Lys Ala Lys Ile Leu Thr Ser Val Ala Leu Leu Ala Cys Ser Gly
  1                      5                      10                      15

Ser Leu Phe Ala Gln Thr Leu Ala Thr Val Asn Gly Gln Lys Ile Asp
  20                      25                      30

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Ser Ser Val Ile Asp Ala Gln Val Ala Ala Phe Arg Ala Glu Asn Ser
 35 40 45
 Arg Ala Glu Asp Thr Pro Gln Leu Arg Gln Ser Leu Leu Glu Asn Glu
 50 55 60
 Val Val Asn Thr Val Val Ala Gln Glu Val Lys Arg Leu Lys Leu Asp
 65 70 75 80
 Arg Ser Ala Glu Phe Lys Asn Ala Leu Ala Lys Leu Arg Ala Glu Ala
 85 90 95
 Lys Lys Ser Gly Asp Asp Lys Lys Pro Ser Phe Lys Thr Val Trp Gln
 100 105 110
 Ala Val Lys Tyr Gly Leu Asn Gly Glu Ala Tyr Ala Leu His Ile Ala
 115 120 125
 Lys Thr Gln Pro Val Ser Glu Gln Glu Val Lys Ala Ala Tyr Asp Asn
 130 135 140
 Ile Ser Gly Phe Tyr Lys Gly Thr Gln Glu Val Gln Leu Gly Glu Ile
 145 150 155 160
 Leu Thr Asp Lys Glu Glu Asn Ala Lys Lys Ala Val Ala Asp Leu Lys
 165 170 175
 Ala Lys Lys Gly Phe Asp Ala Val Leu Lys Gln Tyr Ser Leu Asn Asp
 180 185 190
 Arg Thr Lys Gln Thr Gly Ala Pro Val Gly Tyr Val Pro Leu Lys Asp
 195 200 205
 Leu Glu Gln Gly Val Pro Pro Leu Tyr Gln Ala Ile Lys Asp Leu Lys
 210 215 220
 Lys Gly Glu Phe Thr Ala Thr Pro Leu Lys Asn Gly Asp Phe Tyr Gly
 225 230 235 240
 Val Tyr Tyr Val Asn Asp Ser Arg Glu Val Lys Val Pro Ser Phe Asp
 245 250 255
 Glu Met Lys Gly Gln Ile Ala Gly Asn Leu Gln Ala Glu Arg Ile Asp
 260 265 270
 Arg Ala Val Gly Ala Leu Leu Gly Lys Ala Asn Ile Lys Pro Ala Lys
 275 280 285

<210> 173
 <211> 1794
 <212> DNA
 <213> *Neisseria meningitidis*

<220>
 <221> CDS
 <222> (1)..(1794)

<400> 173
 ttg ccc cga att gcc atg ccc tat ttc gcc ctg ttt gac gat gcc gta 48
 Leu Pro Arg Ile Ala Met Pro Tyr Phe Ala Leu Phe Asp Asp Ala Val
 1 5 10 15
 agc ggt cgc gca aaa cgc tat caa aat cat gtg gaa agc cgt ttt ttc 96
 Ser Gly Arg Ala Lys Arg Tyr Gln Asn His Val Glu Ser Arg Phe Phe
 20 25 30
 cgt ccc gaa gaa ctc gat gct ttg gac ggc gcg ctg caa tcg ggc tgg 144
 Arg Pro Glu Leu Asp Ala Leu Asp Gly Ala Leu Gln Ser Gly Trp
 35 40 45
 caa aaa ggg ctg cat gcc gtg ttg ttt gca gac tac gga ttc ggt ttg 192
 Gln Lys Gly Leu His Ala Val Leu Phe Ala Asp Tyr Gly Phe Gly Leu
 50 55 60
 ccg ctg acg ggg gtc gag tcc gaa cgc ggc ggc aac ctt gcc ctg cac 240
 Pro Leu Thr Gly Val Glu Ser Glu Arg Gly Gly Asn Leu Ala Leu His
 65 70 75 80
 tgg ttt gcc gac tgc gcc gac acc gat gcc gca agc tgg ctt gcc cga 288
 Trp Phe Ala Asp Cys Ala Asp Thr Asp Ala Ala Ser Trp Leu Ala Arg
 85 90 95
 cac tca gac ggc ctc ccc gcc ggc att tcc acg ccg caa tcc tcc gta 336
 His Ser Asp Gly Leu Pro Ala Gly Ile Ser Thr Pro Gln Ser Ser Val
 100 105 110
 tcc gaa gcc gat tac ctc gac cat atc cgc caa atc cac gaa gcc atc 384
 Ser Glu Ala Asp Tyr Leu Asp His Ile Arg Gln Ile His Glu Ala Ile
 115 120 125
 cga cgc ggc gac acc tat caa atc aac tac act acc cgc ctg cac ctg 432
 Arg Arg Gly Asp Thr Tyr Gln Ile Asn Tyr Thr Thr Arg Leu His Leu
 130 135 140

caa gcc tac ggc aat ccc gtc aaa ctt tac caa cgc ctg cgc cag ccc	480
Gln Ala Tyr Gly Asn Pro Val Lys Leu Tyr Gln Arg Leu Arg Gln Pro	
145 150 155 160	
gtc ccc tat gcc gtt ttg tcc cac ctg ccc gat gca caa ggg caa tcc	528
Val Pro Tyr Ala Val Leu Ser His Leu Pro Asp Ala Gln Gly Gln Ser	
165 170 175	
gcg tgg acg ctg tgt ttc tcg ccc gaa ctc ttc ctc aaa atc ggt tcg	576
Ala Trp Thr Leu Cys Phe Ser Pro Glu Leu Phe Leu Lys Ile Gly Ser	
180 185 190	
gac ggc acc atc agc acc gaa ccg atg aaa ggc acc gcg ccg att ttg	624
Asp Gly Thr Ile Ser Thr Glu Pro Met Lys Gly Thr Ala Pro Ile Leu	
195 200 205	
ggc gac gga caa gac gaa cgc cgc gcc gcc gag ttg caa gca gac ccg	672
Gly Asp Gly Gln Asp Glu Arg Arg Ala Ala Glu Leu Gln Ala Asp Pro	
210 215 220	
aaa aac cgc gcc gaa aac gtg atg att gtc gat ttg ctg cgt aac gac	720
Lys Asn Arg Ala Glu Asn Val Met Ile Val Asp Leu Leu Arg Asn Asp	
225 230 235 240	
ctc ggc aaa atc gcc caa acc ggc aca gta tgc gta ccc gaa ccg ttt	768
Leu Gly Lys Ile Ala Gln Thr Gly Thr Val Cys Val Pro Glu Pro Phe	
245 250 255	
aaa gta tcg cgt ttc ggc agc gtt tgg cag atg acc agc acc atc caa	816
Lys Val Ser Arg Phe Gly Ser Val Trp Gln Met Thr Ser Thr Ile Gln	
260 265 270	
gcc caa gcc ttg ccg cac acc tcg ttc gcc gac atc ctc cgc gcc gcc	864
Ala Gln Ala Leu Pro His Thr Ser Phe Ala Asp Ile Leu Arg Ala Ala	
275 280 285	
ttc ccc tgc ggc agc atc acc ggc gcg ccc aaa aaa atg agt atg cag	912
Phe Pro Cys Gly Ser Ile Thr Gly Ala Pro Lys Lys Met Ser Met Gln	
290 295 300	
att atc gaa tcg ctc gaa gcc gaa gcg cgc gga ctt tat acg ggc agc	960
Ile Ile Glu Ser Leu Glu Ala Glu Ala Arg Gly Leu Tyr Thr Gly Ser	
305 310 315 320	
atc ggc tat ttg aac ccg tgt tcc ggc ggc ttg ggg ttt gaa ggc acg	1008
Ile Gly Tyr Leu Asn Pro Cys Ser Gly Gly Leu Gly Phe Glu Gly Thr	
325 330 335	

ttc aac gtc gtt atc cgc acc ttg tgc ctc acg ccg ctt tca gac ggc	1056
Phe Asn Val Val Ile Arg Thr Leu Ser Leu Thr Pro Leu Ser Asp Gly	
340 345 350	
att tat caa ggc gtg tac ggt gtc ggt tcc ggc atc gtc atc gac agc	1104
Ile Tyr Gln Gly Val Tyr Gly Val Gly Ser Gly Ile Val Ile Asp Ser	
355 360 365	
gac cct gcc gcc gaa tat cgg gaa tgc ggc tgg aaa gcc cgt ttc ctc	1152
Asp Pro Ala Ala Glu Tyr Arg Glu Cys Gly Trp Lys Ala Arg Phe Leu	
370 375 380	
aac gaa ttg cgc ccc gac ttc ggc att ttt gaa acc ctg cgc gtg gaa	1200
Asn Glu Leu Arg Pro Asp Phe Gly Ile Phe Glu Thr Leu Arg Val Glu	
385 390 395 400	
aac gga cgc tgc gcc ctg ctc gac cgc cac cta tgc cgt ctg aaa acc	1248
Asn Gly Arg Cys Ala Leu Leu Asp Arg His Leu Cys Arg Leu Lys Thr	
405 410 415	
tcc gcc caa gcc ctc aac ctg ccc ctg ccc gac ggt tgc gaa aat caa	1296
Ser Ala Gln Ala Leu Asn Leu Pro Leu Pro Asp Gly Cys Glu Asn Gln	
420 425 430	
atc aaa caa tac atc gca cgc ttg ccc gac ggc gcg ttc cgc atc aaa	1344
Ile Lys Gln Tyr Ile Ala Arg Leu Pro Asp Gly Ala Phe Arg Ile Lys	
435 440 445	
gcc ctg ctc gct tca gac ggc atc agc ctg tcc cgc gcc gtt tta aac	1392
Ala Leu Leu Ala Ser Asp Gly Ile Ser Leu Ser Arg Ala Val Leu Asn	
450 455 460	
cgt ctg acc gac aaa caa cgc gtc atc att tgc ccg acc atc ctg ccc	1440
Arg Leu Thr Asp Lys Gln Arg Val Ile Ile Ser Pro Thr Ile Leu Pro	
465 470 475 480	
gca caa aac tac ctg cgc cgc ttc aaa acc acc tgc cgc acc gtc ttc	1488
Ala Gln Asn Tyr Leu Arg Arg Phe Lys Thr Thr Cys Arg Thr Val Phe	
485 490 495	
gac caa gcg tgg caa acc gcc gaa aca caa ggc gcg ttc gac agc ctg	1536
Asp Gln Ala Trp Gln Thr Ala Glu Thr Gln Gly Ala Phe Asp Ser Leu	
500 505 510	
ttt ttc aat tca gac ggc atc ctg ctc gaa ggc ggc aga agc aac gtg	1584
Phe Phe Asn Ser Asp Gly Ile Leu Leu Glu Gly Gly Arg Ser Asn Val	
515 520 525	

ttc gtc aaa cat cgc gga caa tgg ctc acg ccc tct tta gat tta gac 1632
 Phe Val Lys His Arg Gly Gln Trp Leu Thr Pro Ser Leu Asp Leu Asp
 530 535 540

att tta aac ggc ata atg cgc caa gcc gtg ttg gac gaa ccg caa aaa 1680
 Ile Leu Asn Gly Ile Met Arg Gln Ala Val Leu Asp Glu Pro Gln Lys
 545 550 555 560

tat ttg caa aca aat caa gta atc gaa aca cac atc aca caa aaa aca 1728
 Tyr Leu Gln Thr Asn Gln Val Ile Glu Thr His Ile Thr Gln Lys Thr
 565 570 575

ctg caa gaa gcc gaa gaa atc cgc ctc tcc aac gcc ttg cgc ggc gta 1776
 Leu Gln Glu Ala Glu Glu Ile Arg Leu Ser Asn Ala Leu Arg Gly Val
 580 585 590

ttt gcc gcc gcc ctt gcc 1794
 Phe Ala Ala Ala Leu Ala
 595

<210> 174

<211> 598

<212> PRT

<213> *Neisseria meningitidis*

<400> 174

Leu Pro Arg Ile Ala Met Pro Tyr Phe Ala Leu Phe Asp Asp Ala Val
 1 5 10 15

Ser Gly Arg Ala Lys Arg Tyr Gln Asn His Val Glu Ser Arg Phe Phe
 20 25 30

Arg Pro Glu Glu Leu Asp Ala Leu Asp Gly Ala Leu Gln Ser Gly Trp
 35 40 45

Gln Lys Gly Leu His Ala Val Leu Phe Ala Asp Tyr Gly Phe Gly Leu
 50 55 60

Pro Leu Thr Gly Val Glu Ser Glu Arg Gly Gly Asn Leu Ala Leu His
 65 70 75 80

Trp Phe Ala Asp Cys Ala Asp Thr Asp Ala Ala Ser Trp Leu Ala Arg
 85 90 95

His Ser Asp Gly Leu Pro Ala Gly Ile Ser Thr Pro Gln Ser Ser Val
 100 105 110

Ser Glu Ala Asp Tyr Leu Asp His Ile Arg Gln Ile His Glu Ala Ile
 115 120 125
 Arg Arg Gly Asp Thr Tyr Gln Ile Asn Tyr Thr Thr Arg Leu His Leu
 130 135 140
 Gln Ala Tyr Gly Asn Pro Val Lys Leu Tyr Gln Arg Leu Arg Gln Pro
 145 150 155 160
 Val Pro Tyr Ala Val Leu Ser His Leu Pro Asp Ala Gln Gly Gln Ser
 165 170 175
 Ala Trp Thr Leu Cys Phe Ser Pro Glu Leu Phe Leu Lys Ile Gly Ser
 180 185 190
 Asp Gly Thr Ile Ser Thr Glu Pro Met Lys Gly Thr Ala Pro Ile Leu
 195 200 205
 Gly Asp Gly Gln Asp Glu Arg Arg Ala Ala Glu Leu Gln Ala Asp Pro
 210 215 220
 Lys Asn Arg Ala Glu Asn Val Met Ile Val Asp Leu Leu Arg Asn Asp
 225 230 235 240
 Leu Gly Lys Ile Ala Gln Thr Gly Thr Val Cys Val Pro Glu Pro Phe
 245 250 255
 Lys Val Ser Arg Phe Gly Ser Val Trp Gln Met Thr Ser Thr Ile Gln
 260 265 270
 Ala Gln Ala Leu Pro His Thr Ser Phe Ala Asp Ile Leu Arg Ala Ala
 275 280 285
 Phe Pro Cys Gly Ser Ile Thr Gly Ala Pro Lys Lys Met Ser Met Gln
 290 295 300
 Ile Ile Glu Ser Leu Glu Ala Glu Ala Arg Gly Leu Tyr Thr Gly Ser
 305 310 315 320
 Ile Gly Tyr Leu Asn Pro Cys Ser Gly Gly Leu Gly Phe Glu Gly Thr
 325 330 335
 Phe Asn Val Val Ile Arg Thr Leu Ser Leu Thr Pro Leu Ser Asp Gly
 340 345 350
 Ile Tyr Gln Gly Val Tyr Gly Val Gly Ser Gly Ile Val Ile Asp Ser
 355 360 365

Asp Pro Ala Ala Glu Tyr Arg Glu Cys Gly Trp Lys Ala Arg Phe Leu
 370 375 380

Asn Glu Leu Arg Pro Asp Phe Gly Ile Phe Glu Thr Leu Arg Val Glu
 385 390 395 400

Asn Gly Arg Cys Ala Leu Leu Asp Arg His Leu Cys Arg Leu Lys Thr
 405 410 415

Ser Ala Gln Ala Leu Asn Leu Pro Leu Pro Asp Gly Cys Glu Asn Gln
 420 425 430

Ile Lys Gln Tyr Ile Ala Arg Leu Pro Asp Gly Ala Phe Arg Ile Lys
 435 440 445

Ala Leu Leu Ala Ser Asp Gly Ile Ser Leu Ser Arg Ala Val Leu Asn
 450 455 460

Arg Leu Thr Asp Lys Gln Arg Val Ile Ile Ser Pro Thr Ile Leu Pro
 465 470 475 480

Ala Gln Asn Tyr Leu Arg Arg Phe Lys Thr Thr Cys Arg Thr Val Phe
 485 490 495

Asp Gln Ala Trp Gln Thr Ala Glu Thr Gln Gly Ala Phe Asp Ser Leu
 500 505 510

Phe Phe Asn Ser Asp Gly Ile Leu Leu Glu Gly Gly Arg Ser Asn Val
 515 520 525

Phe Val Lys His Arg Gly Gln Trp Leu Thr Pro Ser Leu Asp Leu Asp
 530 535 540

Ile Leu Asn Gly Ile Met Arg Gln Ala Val Leu Asp Glu Pro Gln Lys
 545 550 555 560

Tyr Leu Gln Thr Asn Gln Val Ile Glu Thr His Ile Thr Gln Lys Thr
 565 570 575

Leu Gln Glu Ala Glu Glu Ile Arg Leu Ser Asn Ala Leu Arg Gly Val
 580 585 590

Phe Ala Ala Ala Leu Ala
 595

<210> 175

<211> 279

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(279)

<400> 175

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ttg caa aac ttt cgg aaa ccc aat atg cag acc gtt acc atg tac aca 48
Leu Gln Asn Phe Arg Lys Pro Asn Met Gln Thr Val Thr Met Tyr Thr
  1             5             10            15

```

```

ggt ccg ttt tgc ccc tac tgc acg atg gcg aaa agg ctg ctg cac gcg 96
Gly Pro Phe Cys Pro Tyr Cys Thr Met Ala Lys Arg Leu Leu His Ala
          20             25            30

```

```

gca ggt gtc gga cat atc gac gaa atc cgt gtc gat gca agt ccc gaa 144
Ala Gly Val Gly His Ile Asp Glu Ile Arg Val Asp Ala Ser Pro Glu
          35             40            45

```

```

gcc ttt gcc gaa atg cag cag ctt tcc gga cag cgc agc gtg ccg cag 192
Ala Phe Ala Glu Met Gln Gln Leu Ser Gly Gln Arg Ser Val Pro Gln
          50             55            60

```

```

att ttc atc ggc gaa acg cac gtc ggc gga ttt acc gac ctc tac cgc 240
Ile Phe Ile Gly Glu Thr His Val Gly Gly Phe Thr Asp Leu Tyr Arg
          65             70            75            80

```

```

ctc cag cag gaa ggc ggg ctg gac gga ctg ctg aac cct 279
Leu Gln Gln Glu Gly Gly Leu Asp Gly Leu Leu Asn Pro
          85             90

```

<210> 176

<211> 93

<212> PRT

<213> *Neisseria meningitidis*

<400> 176

```

Leu Gln Asn Phe Arg Lys Pro Asn Met Gln Thr Val Thr Met Tyr Thr
  1             5             10            15

```

```

Gly Pro Phe Cys Pro Tyr Cys Thr Met Ala Lys Arg Leu Leu His Ala
          20             25            30

```

```

Ala Gly Val Gly His Ile Asp Glu Ile Arg Val Asp Ala Ser Pro Glu

```


35 40 45
 Ala Phe Ala Glu Met Gln Gln Leu Ser Gly Gln Arg Ser Val Pro Gln
 50 55 60
 Ile Phe Ile Gly Glu Thr His Val Gly Gly Phe Thr Asp Leu Tyr Arg
 65 70 75 80
 Leu Gln Gln Glu Gly Gly Leu Asp Gly Leu Leu Asn Pro
 85 90

 <210> 177
 <211> 2274
 <212> DNA
 <213> Neisseria meningitidis

 <220>
 <221> CDS
 <222> (1)..(2274)

 <400> 177
 atg aca aca tta cat ttc tca ggc ttc ccg cgt gtc ggt gcc ttc cgc 48
 Met Thr Thr Leu His Phe Ser Gly Phe Pro Arg Val Gly Ala Phe Arg
 1 5 10 15

 gaa ttg aaa ttc gca caa gaa aaa tac tgg cgc aaa gaa atc agc gag 96
 Glu Leu Lys Phe Ala Gln Glu Lys Tyr Trp Arg Lys Glu Ile Ser Glu
 20 25 30

 caa gaa ttg ctg gct gtt gct aaa gac ttg cgc gag aaa aac tgg aaa 144
 Gln Glu Leu Leu Ala Val Ala Lys Asp Leu Arg Glu Lys Asn Trp Lys
 35 40 45

 cac cag gcc gct gcc aac gcc gat tac gtt gcc gta ggc gat ttc act 192
 His Gln Ala Ala Ala Asn Ala Asp Tyr Val Ala Val Gly Asp Phe Thr
 50 55 60

 ttc tac gac cac atc ctc gac ctg caa gtc gcc acc ggc gcg att ccc 240
 Phe Tyr Asp His Ile Leu Asp Leu Gln Val Ala Thr Gly Ala Ile Pro
 65 70 75 80

 gcc cgc ttc ggc ttc gac agc caa aac cta tct ttg gaa caa ttc ttc 288
 Ala Arg Phe Gly Phe Asp Ser Gln Asn Leu Ser Leu Glu Gln Phe Phe
 85 90 95

 caa ctg gcg cgc ggt aac aaa gac caa ttc gct atc gaa atg acc aaa 336

344

345

Val Leu Val His Gly Glu Ala Glu Arg Asn Asp Met Val Glu Tyr Phe	
485 490 495	
ggc gaa ttg ttg agc ggt ttt gca ttc acc caa tat ggc tgg gta caa	1536
Gly Glu Leu Leu Ser Gly Phe Ala Phe Thr Gln Tyr Gly Trp Val Gln	
500 505 510	
agc tac ggt tct cgc tgc gtt aaa cca ccg att atc ttc ggt gac gta	1584
Ser Tyr Gly Ser Arg Cys Val Lys Pro Pro Ile Ile Phe Gly Asp Val	
515 520 525	
agc cgt cct gaa gcc atg acc gta gct tgg tct act tac gca caa agc	1632
Ser Arg Pro Glu Ala Met Thr Val Ala Trp Ser Thr Tyr Ala Gln Ser	
530 535 540	
ctg acc aaa cgc ccg atg aaa ggt atg ttg acc ggc cct gta acc att	1680
Leu Thr Lys Arg Pro Met Lys Gly Met Leu Thr Gly Pro Val Thr Ile	
545 550 555 560	
ctg caa tgg tct ttc gtc cgc aac gat att cct cgc tct acc gtg tgt	1728
Leu Gln Trp Ser Phe Val Arg Asn Asp Ile Pro Arg Ser Thr Val Cys	
565 570 575	
aaa caa atc gca ctg gct ctg aac gac gaa gta ttg gat ctg gaa aaa	1776
Lys Gln Ile Ala Leu Ala Leu Asn Asp Glu Val Leu Asp Leu Glu Lys	
580 585 590	
gca ggc atc aaa gtc atc caa atc gac gaa cct gcc atc cgc gaa ggc	1824
Ala Gly Ile Lys Val Ile Gln Ile Asp Glu Pro Ala Ile Arg Glu Gly	
595 600 605	
ctg cct ttg aaa cgt gcc gat tgg gat gcc tac ctg aac tgg gcc ggc	1872
Leu Pro Leu Lys Arg Ala Asp Trp Asp Ala Tyr Leu Asn Trp Ala Gly	
610 615 620	
gaa tct ttc cgc ctg tcc tct acc ggt tgc gaa gac agc act caa atc	1920
Glu Ser Phe Arg Leu Ser Ser Thr Gly Cys Glu Asp Ser Thr Gln Ile	
625 630 635 640	
cac acc cat atg tgc tac tct gag ttc aac gac atc ctg cct gcg att	1968
His Thr His Met Cys Tyr Ser Glu Phe Asn Asp Ile Leu Pro Ala Ile	
645 650 655	
gct gca atg gat gcg gac gtg atc acc atc gag act tca cgt tcc gac	2016
Ala Ala Met Asp Ala Asp Val Ile Thr Ile Glu Thr Ser Arg Ser Asp	
660 665 670	
atg gaa ctc ttg acc gcg ttc ggc gaa ttc aaa tac ccg aac gac atc	2064

Met Glu Leu Leu Thr Ala Phe Gly Glu Phe Lys Tyr Pro Asn Asp Ile
675 680 685

ggc ccg ggg gtt tac gac atc cac agc ccg cgc gta ccg aca gaa gcc 2112
Gly Pro Gly Val Tyr Asp Ile His Ser Pro Arg Val Pro Thr Glu Ala
690 695 700

gaa gtg gag cac ctg ttg cgc aaa gcc atc gag gtt gta ccg gtt gaa 2160
Glu Val Glu His Leu Leu Arg Lys Ala Ile Glu Val Val Pro Val Glu
705 710 715 720

cgt ctg tgg gtt aac ccg gac tgc ggc ctg aaa aca cgc ggc tgg aaa 2208
Arg Leu Trp Val Asn Pro Asp Cys Gly Leu Lys Thr Arg Gly Trp Lys
725 730 735

gaa act ctg gaa caa ctc caa gtg atg atg aac gta acc cac aaa ttg 2256
Glu Thr Leu Glu Gln Leu Gln Val Met Met Asn Val Thr His Lys Leu
740 745 750

cgt gcc gaa ttg gcg aaa 2274
Arg Ala Glu Leu Ala Lys
755

<210> 178

<211> 758

<212> PRT

<213> Neisseria meningitidis

<400> 178

Met Thr Thr Leu His Phe Ser Gly Phe Pro Arg Val Gly Ala Phe Arg
1 5 10 15

Glu Leu Lys Phe Ala Gln Glu Lys Tyr Trp Arg Lys Glu Ile Ser Glu
20 25 30

Gln Glu Leu Leu Ala Val Ala Lys Asp Leu Arg Glu Lys Asn Trp Lys
35 40 45

His Gln Ala Ala Ala Asn Ala Asp Tyr Val Ala Val Gly Asp Phe Thr
50 55 60

Phe Tyr Asp His Ile Leu Asp Leu Gln Val Ala Thr Gly Ala Ile Pro
65 70 75 80

Ala Arg Phe Gly Phe Asp Ser Gln Asn Leu Ser Leu Glu Gln Phe Phe
85 90 95

Gln Leu Ala Arg Gly Asn Lys Asp Gln Phe Ala Ile Glu Met Thr Lys
 100 105 110

Trp Phe Asp Thr Asn Tyr His Tyr Leu Val Pro Glu Phe His Ala Asp
 115 120 125

Thr Glu Phe Lys Ala Asn Ala Lys His Tyr Val Gln Gln Leu Gln Glu
 130 135 140

Ala Gln Ala Leu Gly Leu Lys Ala Lys Pro Thr Val Val Gly Pro Leu
 145 150 155 160

Thr Phe Leu Trp Val Gly Lys Glu Lys Gly Ala Val Glu Phe Asp Arg
 165 170 175

Leu Ser Leu Leu Pro Lys Leu Leu Pro Val Tyr Val Glu Ile Leu Thr
 180 185 190

Ala Leu Val Glu Ala Gly Ala Glu Trp Ile Gln Ile Asp Glu Pro Ala
 195 200 205

Leu Thr Val Asp Leu Pro Lys Glu Trp Val Glu Ala Tyr Lys Asp Val
 210 215 220

Tyr Ala Thr Leu Ser Lys Val Ser Ala Lys Ile Leu Leu Ser Thr Tyr
 225 230 235 240

Phe Gly Ser Val Ala Glu His Ala Ala Leu Leu Lys Ser Leu Pro Val
 245 250 255

Asp Gly Leu His Ile Asp Leu Val Arg Ala Pro Glu Gln Leu Asp Ala
 260 265 270

Phe Ala Asp Tyr Asp Lys Val Leu Ser Ala Gly Val Ile Asp Gly Arg
 275 280 285

Asn Ile Trp Arg Ala Asn Leu Asn Lys Val Leu Glu Thr Val Glu Leu
 290 295 300

Leu Gln Ala Lys Leu Gly Asp Arg Leu Trp Ile Ser Ser Ser Cys Ser
 305 310 315 320

Leu Leu His Thr Pro Phe Asp Leu Ser Val Glu Glu Lys Leu Lys Ala
 325 330 335

Asn Lys Pro Asp Leu Tyr Ser Trp Leu Ala Phe Thr Leu Gln Lys Thr
 340 345 350

Gln Glu Leu Arg Val Leu Lys Ala Ala Leu Asn Glu Gly Arg Asp Ser
 355 360 365

Val Ala Glu Glu Leu Ala Ala Ser Gln Ala Ala Ala Asp Ser Arg Ala
 370 375 380

Asn Ser Ser Glu Ile His Arg Ala Asp Val Ala Lys Arg Leu Ala Asp
 385 390 395 400

Leu Pro Ala Asn Ala Asp Gln Arg Lys Ser Pro Phe Ala Asp Arg Ile
 405 410 415

Lys Ala Gln Gln Ala Trp Leu Asn Leu Pro Leu Leu Pro Thr Thr Asn
 420 425 430

Ile Gly Ser Phe Pro Gln Thr Thr Glu Ile Arg Gln Ala Arg Ala Ala
 435 440 445

Phe Lys Lys Gly Glu Leu Ser Ala Ala Asp Tyr Glu Ala Ala Met Lys
 450 455 460

Lys Glu Ile Ala Leu Val Val Glu Glu Gln Glu Lys Leu Asp Leu Asp
 465 470 475 480

Val Leu Val His Gly Glu Ala Glu Arg Asn Asp Met Val Glu Tyr Phe
 485 490 495

Gly Glu Leu Leu Ser Gly Phe Ala Phe Thr Gln Tyr Gly Trp Val Gln
 500 505 510

Ser Tyr Gly Ser Arg Cys Val Lys Pro Pro Ile Ile Phe Gly Asp Val
 515 520 525

Ser Arg Pro Glu Ala Met Thr Val Ala Trp Ser Thr Tyr Ala Gln Ser
 530 535 540

Leu Thr Lys Arg Pro Met Lys Gly Met Leu Thr Gly Pro Val Thr Ile
 545 550 555 560

Leu Gln Trp Ser Phe Val Arg Asn Asp Ile Pro Arg Ser Thr Val Cys
 565 570 575

Lys Gln Ile Ala Leu Ala Leu Asn Asp Glu Val Leu Asp Leu Glu Lys
 580 585 590

Ala Gly Ile Lys Val Ile Gln Ile Asp Glu Pro Ala Ile Arg Glu Gly
 595 600 605

Leu Pro Leu Lys Arg Ala Asp Trp Asp Ala Tyr Leu Asn Trp Ala Gly
 610 615 620
 Glu Ser Phe Arg Leu Ser Ser Thr Gly Cys Glu Asp Ser Thr Gln Ile
 625 630 635 640
 His Thr His Met Cys Tyr Ser Glu Phe Asn Asp Ile Leu Pro Ala Ile
 645 650 655
 Ala Ala Met Asp Ala Asp Val Ile Thr Ile Glu Thr Ser Arg Ser Asp
 660 665 670
 Met Glu Leu Leu Thr Ala Phe Gly Glu Phe Lys Tyr Pro Asn Asp Ile
 675 680 685
 Gly Pro Gly Val Tyr Asp Ile His Ser Pro Arg Val Pro Thr Glu Ala
 690 695 700
 Glu Val Glu His Leu Leu Arg Lys Ala Ile Glu Val Val Pro Val Glu
 705 710 715 720
 Arg Leu Trp Val Asn Pro Asp Cys Gly Leu Lys Thr Arg Gly Trp Lys
 725 730 735
 Glu Thr Leu Glu Gln Leu Gln Val Met Met Asn Val Thr His Lys Leu
 740 745 750
 Arg Ala Glu Leu Ala Lys
 755

<210> 179

<211> 867

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(867)

<400> 179

atg aaa ccc ata cgg aaa gcc gtc ttc ccc gtc gca ggg atg gga aca 48
 Met Lys Pro Ile Arg Lys Ala Val Phe Pro Val Ala Gly Met Gly Thr
 1 5 10 15
 cgc ttc ctg ccc gcc acc aag gca agc ccg aaa gaa atg ctg ccc atc 96
 Arg Phe Leu Pro Ala Thr Lys Ala Ser Pro Lys Glu Met Leu Pro Ile

20	25	30	
gtc gac aag ccg ctg atc caa tac gcc gta gaa gaa gcc gtg gaa gcc			144
Val Asp Lys Pro Leu Ile Gln Tyr Ala Val Glu Glu Ala Val Glu Ala			
35	40	45	
ggc tgc acg gaa atg gtg ttt gtt acc gga cgc aac aaa cgc agc atc			192
Gly Cys Thr Glu Met Val Phe Val Thr Gly Arg Asn Lys Arg Ser Ile			
50	55	60	
gaa gac cat ttc gac aag gca tac gaa ctc gaa acc gag ttg gaa atg			240
Glu Asp His Phe Asp Lys Ala Tyr Glu Leu Glu Thr Glu Leu Glu Met			
65	70	75	80
cgc cat aaa gac aaa ttg ttg gaa cac gtc cgc aac atc ctg ccg ccg			288
Arg His Lys Asp Lys Leu Leu Glu His Val Arg Asn Ile Leu Pro Pro			
85	90	95	
aac att acc tgc ctc tac atc cgt cag gcg gaa gca ctg ggc ttg gga			336
Asn Ile Thr Cys Leu Tyr Ile Arg Gln Ala Glu Ala Leu Gly Leu Gly			
100	105	110	
cac gcc gtc ttg tgc gcc cgc gcc gcc atc ggc gac gaa ccc ttt gcc			384
His Ala Val Leu Cys Ala Arg Ala Ala Ile Gly Asp Glu Pro Phe Ala			
115	120	125	
gtg att ttg gct gac gat tta atc gat gcc caa aaa ggc gcg ctc aaa			432
Val Ile Leu Ala Asp Asp Leu Ile Asp Ala Gln Lys Gly Ala Leu Lys			
130	135	140	
caa atg gtc gaa gtg tac gaa cgc agc ggc aac agc att ttg ggc gta			480
Gln Met Val Glu Val Tyr Glu Arg Ser Gly Asn Ser Ile Leu Gly Val			
145	150	155	160
gaa act gtc gaa ccg tcg caa acc ggc tca tac ggc atc gtc gaa acc			528
Glu Thr Val Glu Pro Ser Gln Thr Gly Ser Tyr Gly Ile Val Glu Thr			
165	170	175	
gaa cag ctc aaa cag ttc caa cgc att acc ggc att gtc gaa aaa ccc			576
Glu Gln Leu Lys Gln Phe Gln Arg Ile Thr Gly Ile Val Glu Lys Pro			
180	185	190	
aag ccc gaa gac gcg ccc tcc aac ctt gcc gtc gtc gga cgc tac atc			624
Lys Pro Glu Asp Ala Pro Ser Asn Leu Ala Val Val Gly Arg Tyr Ile			
195	200	205	
ctt acc ccg cgc att ttc gac tta ctg acc gga ctg ccg cgc ggc gcg			672
Leu Thr Pro Arg Ile Phe Asp Leu Leu Thr Gly Leu Pro Arg Gly Ala			

210 215 220
 ggc aac gaa atc cag ctt aca gac ggc atc gcc aag ctg ctc gat cac 720
 Gly Asn Glu Ile Gln Leu Thr Asp Gly Ile Ala Lys Leu Leu Asp His
 225 230 235 240
 gaa ttt gtc cta gcg cac ccc ttt gaa ggc acg cgc tac gac tgc ggc 768
 Glu Phe Val Leu Ala His Pro Phe Glu Gly Thr Arg Tyr Asp Cys Gly
 245 250 255
 agc aaa ctg ggc tat ctg gaa gcc acc gtc gcc tac ggc ctg aaa cac 816
 Ser Lys Leu Gly Tyr Leu Glu Ala Thr Val Ala Tyr Gly Leu Lys His
 260 265 270
 ccc gaa acc ggc gaa ccc ttc cgc cgc ctt ttg gaa aaa tac cgt acc 864
 Pro Glu Thr Gly Glu Pro Phe Arg Arg Leu Leu Glu Lys Tyr Arg Thr
 275 280 285
 gaa 867
 Glu

<210> 180

<211> 289

<212> PRT

<213> Neisseria meningitidis

<400> 180

Met Lys Pro Ile Arg Lys Ala Val Phe Pro Val Ala Gly Met Gly Thr
 1 5 10 15

Arg Phe Leu Pro Ala Thr Lys Ala Ser Pro Lys Glu Met Leu Pro Ile
 20 25 30

Val Asp Lys Pro Leu Ile Gln Tyr Ala Val Glu Glu Ala Val Glu Ala
 35 40 45

Gly Cys Thr Glu Met Val Phe Val Thr Gly Arg Asn Lys Arg Ser Ile
 50 55 60

Glu Asp His Phe Asp Lys Ala Tyr Glu Leu Glu Thr Glu Leu Glu Met
 65 70 75 80

Arg His Lys Asp Lys Leu Leu Glu His Val Arg Asn Ile Leu Pro Pro
 85 90 95

Asn Ile Thr Cys Leu Tyr Ile Arg Gln Ala Glu Ala Leu Gly Leu Gly
 100 105 110

```

His Ala Val Leu Cys Ala Arg Ala Ala Ile Gly Asp Glu Pro Phe Ala
      115              120              125

Val Ile Leu Ala Asp Asp Leu Ile Asp Ala Gln Lys Gly Ala Leu Lys
      130              135              140

Gln Met Val Glu Val Tyr Glu Arg Ser Gly Asn Ser Ile Leu Gly Val
      145              150              155              160

Glu Thr Val Glu Pro Ser Gln Thr Gly Ser Tyr Gly Ile Val Glu Thr
      165              170              175

Glu Gln Leu Lys Gln Phe Gln Arg Ile Thr Gly Ile Val Glu Lys Pro
      180              185              190

Lys Pro Glu Asp Ala Pro Ser Asn Leu Ala Val Val Gly Arg Tyr Ile
      195              200              205

Leu Thr Pro Arg Ile Phe Asp Leu Leu Thr Gly Leu Pro Arg Gly Ala
      210              215              220

Gly Asn Glu Ile Gln Leu Thr Asp Gly Ile Ala Lys Leu Leu Asp His
      225              230              235              240

Glu Phe Val Leu Ala His Pro Phe Glu Gly Thr Arg Tyr Asp Cys Gly
      245              250              255

Ser Lys Leu Gly Tyr Leu Glu Ala Thr Val Ala Tyr Gly Leu Lys His
      260              265              270

Pro Glu Thr Gly Glu Pro Phe Arg Arg Leu Leu Glu Lys Tyr Arg Thr
      275              280              285

Glu

```

<210> 181

<211> 603

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(603)

<400> 181

atg aca ata atg aat atc aat act tcc gaa aat aaa gat gct gtt gcc	48
Met Thr Ile Met Asn Ile Asn Thr Ser Glu Asn Lys Asp Ala Val Ala	
1 5 10 15	
 gaa cac acc gga caa tgg ttg gaa aaa gcc gtc atc ggt ctg aac ctg	96
Glu His Thr Gly Gln Trp Leu Glu Lys Ala Val Ile Gly Leu Asn Leu	
20 25 30	
 tgt ccc ttt gcc aaa gcc ccc cac gtt aaa aac ctt gtc cgc atc gca	144
Cys Pro Phe Ala Lys Ala Pro His Val Lys Asn Leu Val Arg Ile Ala	
35 40 45	
 atc agc gaa gcc aaa cac ctt gac agt ttt ttg gaa gac ttg gac gaa	192
Ile Ser Glu Ala Lys His Leu Asp Ser Phe Leu Glu Asp Leu Asp Glu	
50 55 60	
 gaa ctg cag cga ctg ggc aat aca ccc gcc acc gaa ctg gaa act acc	240
Glu Leu Gln Arg Leu Gly Asn Thr Pro Ala Thr Glu Leu Glu Thr Thr	
65 70 75 80	
 ctg ctg gtt cac ccg acc cta ttc ccc gat ttc gac gta ttc aac gat	288
Leu Leu Val His Pro Thr Leu Phe Pro Asp Phe Asp Val Phe Asn Asp	
85 90 95	
 atg ctc gac att gcc gat gcc gcc gtt gtc gaa aac ggc ttg gaa ggc	336
Met Leu Asp Ile Ala Asp Ala Ala Val Val Glu Asn Gly Leu Glu Gly	
100 105 110	
 atc atc caa atc gcc ccg ttt cat ccc gat ttc caa ttt gaa ggc acg	384
Ile Ile Gln Ile Ala Pro Phe His Pro Asp Phe Gln Phe Glu Gly Thr	
115 120 125	
 gat tca gac gac atc ggc aac tac acc aac cgt tct ccc tat ccg acg	432
Asp Ser Asp Asp Ile Gly Asn Tyr Thr Asn Arg Ser Pro Tyr Pro Thr	
130 135 140	
 ctg cac ctc atc cgc gaa gac agc att gcc aaa gcc gca caa gcc ttt	480
Leu His Leu Ile Arg Glu Asp Ser Ile Ala Lys Ala Ala Gln Ala Phe	
145 150 155 160	
 ccc gac gct tcg gca ata ttc gaa cgc aat atc gcc ctg ctg gaa aaa	528
Pro Asp Ala Ser Ala Ile Phe Glu Arg Asn Ile Ala Leu Leu Glu Lys	
165 170 175	
 atg gga cat gag ggc tgg gca aaa ctc ggt atc aca tcc tgc cct tat	576
Met Gly His Glu Gly Trp Ala Lys Leu Gly Ile Thr Ser Cys Pro Tyr	
180 185 190	

ccg cac aat aag aaa aat att tca aaa
 Pro His Asn Lys Lys Asn Ile Ser Lys
 195 200

603

<210> 182

<211> 201

<212> PRT

<213> Neisseria meningitidis

<400> 182

Met Thr Ile Met Asn Ile Asn Thr Ser Glu Asn Lys Asp Ala Val Ala
 1 5 10 15

Glu His Thr Gly Gln Trp Leu Glu Lys Ala Val Ile Gly Leu Asn Leu
 20 25 30

Cys Pro Phe Ala Lys Ala Pro His Val Lys Asn Leu Val Arg Ile Ala
 35 40 45

Ile Ser Glu Ala Lys His Leu Asp Ser Phe Leu Glu Asp Leu Asp Glu
 50 55 60

Glu Leu Gln Arg Leu Gly Asn Thr Pro Ala Thr Glu Leu Glu Thr Thr
 65 70 75 80

Leu Leu Val His Pro Thr Leu Phe Pro Asp Phe Asp Val Phe Asn Asp
 85 90 95

Met Leu Asp Ile Ala Asp Ala Ala Val Val Glu Asn Gly Leu Glu Gly
 100 105 110

Ile Ile Gln Ile Ala Pro Phe His Pro Asp Phe Gln Phe Glu Gly Thr
 115 120 125

Asp Ser Asp Asp Ile Gly Asn Tyr Thr Asn Arg Ser Pro Tyr Pro Thr
 130 135 140

Leu His Leu Ile Arg Glu Asp Ser Ile Ala Lys Ala Ala Gln Ala Phe
 145 150 155 160

Pro Asp Ala Ser Ala Ile Phe Glu Arg Asn Ile Ala Leu Leu Glu Lys
 165 170 175

Met Gly His Glu Gly Trp Ala Lys Leu Gly Ile Thr Ser Cys Pro Tyr
 180 185 190

Pro His Asn Lys Lys Asn Ile Ser Lys
 195 200

<210> 183

<211> 1338

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1338)

<400> 183

ttg cac acc ctt cta gct ttt atc ttc gcc atc ctg att ttg gtc agc 48
 Leu His Thr Leu Leu Ala Phe Ile Phe Ala Ile Leu Ile Leu Val Ser
 1 5 10 15

ctg cac gaa ttc gga cac tac atc gtc gcc aga ttg tgc ggc gtc aag 96
 Leu His Glu Phe Gly His Tyr Ile Val Ala Arg Leu Cys Gly Val Lys
 20 25 30

gtt gtg cgt ttt tcc gtc ggc ttc ggc aaa ccg ttt ttc acc cga aag 144
 Val Val Arg Phe Ser Val Gly Phe Gly Lys Pro Phe Phe Thr Arg Lys
 35 40 45

cgc ggc gac acc gaa tgg tgc ctc gcc ccg att ccg ttg ggc ggt tac 192
 Arg Gly Asp Thr Glu Trp Cys Leu Ala Pro Ile Pro Leu Gly Gly Tyr
 50 55 60

gtc aaa atg gtc gac acg cgc gaa ggc gaa gta tca gaa gcc gat tta 240
 Val Lys Met Val Asp Thr Arg Glu Gly Glu Val Ser Glu Ala Asp Leu
 65 70 75 80

ccc tac gct ttt gac aaa caa cac ccc gcc aag cgc atc gcc atc gtc 288
 Pro Tyr Ala Phe Asp Lys Gln His Pro Ala Lys Arg Ile Ala Ile Val
 85 90 95

gcc gcc ggc ccg ctg acc aac ctc gca ctg gcg gtt ttg ctg tac gga 336
 Ala Ala Gly Pro Leu Thr Asn Leu Ala Leu Ala Val Leu Leu Tyr Gly
 100 105 110

ctg agc ttt tcc ttc ggc gtt acc gaa ctg cgc ccc tat gtc ggc aca 384
 Leu Ser Phe Ser Phe Gly Val Thr Glu Leu Arg Pro Tyr Val Gly Thr
 115 120 125

gtc gaa ccc gac acc att gcc gcc cgc gcc ggc ttc caa agc ggc gac 432

357

Phe Gly Met Gly Trp Glu Lys Thr Val Ser His Ser Trp Thr Thr Leu
 325 330 335

 aaa ttt ttc ggc aaa cta atc agc ggc aac gcc tcc gtc agc cat att 1056
 Lys Phe Phe Gly Lys Leu Ile Ser Gly Asn Ala Ser Val Ser His Ile
 340 345 350

 tcc ggt ccg ctg acc att gcc gat att gcc gga cag tcc gcc gaa ctc 1104
 Ser Gly Pro Leu Thr Ile Ala Asp Ile Ala Gly Gln Ser Ala Glu Leu
 355 360 365

 ggc ttg caa agt tat ttg gaa ttt ttg gca ctg gtc agc atc agc ctc 1152
 Gly Leu Gln Ser Tyr Leu Glu Phe Leu Ala Leu Val Ser Ile Ser Leu
 370 375 380

 ggc gtg ctg aac ctg ctg ccc gtc ccc gtt ttg gac ggc ggc cac ctc 1200
 Gly Val Leu Asn Leu Leu Pro Val Pro Val Leu Asp Gly Gly His Leu
 385 390 395 400

 gtg ttt tat act gcc gaa tgg ata cgc ggc aaa cct ttg ggc gaa cgc 1248
 Val Phe Tyr Thr Ala Glu Trp Ile Arg Gly Lys Pro Leu Gly Glu Arg
 405 410 415

 gtc caa aac atc ggt ttg cgc ttc ggg ctt gcc ctc atg atg ctg atg 1296
 Val Gln Asn Ile Gly Leu Arg Phe Gly Leu Ala Leu Met Met Leu Met
 420 425 430

 atg gcg gtc gcc ttc ttc aac gac gtt acc cgg ctg ctc ggt 1338
 Met Ala Val Ala Phe Phe Asn Asp Val Thr Arg Leu Leu Gly
 435 440 445

<210> 184

<211> 446

<212> PRT

<213> Neisseria meningitidis

<400> 184

Leu His Thr Leu Leu Ala Phe Ile Phe Ala Ile Leu Ile Leu Val Ser
 1 5 10 15

Leu His Glu Phe Gly His Tyr Ile Val Ala Arg Leu Cys Gly Val Lys
 20 25 30

Val Val Arg Phe Ser Val Gly Phe Gly Lys Pro Phe Phe Thr Arg Lys
 35 40 45

Arg Gly Asp Thr Glu Trp Cys Leu Ala Pro Ile Pro Leu Gly Gly Tyr

50	55	60
Val Lys Met Val Asp Thr Arg Glu Gly Glu Val Ser Glu Ala Asp Leu		
65	70	75 80
Pro Tyr Ala Phe Asp Lys Gln His Pro Ala Lys Arg Ile Ala Ile Val		
	85	90 95
Ala Ala Gly Pro Leu Thr Asn Leu Ala Leu Ala Val Leu Leu Tyr Gly		
	100	105 110
Leu Ser Phe Ser Phe Gly Val Thr Glu Leu Arg Pro Tyr Val Gly Thr		
	115	120 125
Val Glu Pro Asp Thr Ile Ala Ala Arg Ala Gly Phe Gln Ser Gly Asp		
	130	135 140
Lys Ile Gln Ser Val Asn Gly Thr Pro Val Ala Asp Trp Gly Ser Ala		
145	150	155 160
Gln Thr Glu Ile Val Leu Asn Leu Glu Ala Gly Lys Val Ala Val Gly		
	165	170 175
Val Gln Thr Ala Ser Gly Ala Gln Thr Val Arg Thr Ile Asp Ala Ala		
	180	185 190
Gly Thr Pro Glu Ala Gly Lys Ile Ala Lys Asn Gln Gly Tyr Ile Gly		
	195	200 205
Leu Met Pro Phe Lys Ile Thr Thr Val Ala Gly Gly Val Glu Lys Gly		
	210	215 220
Ser Pro Ala Glu Lys Ala Gly Leu Lys Pro Gly Asp Arg Leu Thr Ala		
225	230	235 240
Ala Asp Gly Lys Pro Ile Ala Ser Trp Gln Glu Trp Ala Asn Leu Thr		
	245	250 255
Arg Gln Ser Pro Gly Lys Lys Ile Thr Leu Thr Tyr Glu Arg Ala Gly		
	260	265 270
Gln Thr His Thr Ala Asp Ile Arg Pro Asp Thr Val Glu Gln Pro Asp		
	275	280 285
His Thr Leu Ile Gly Arg Val Gly Leu Arg Pro Gln Pro Asp Arg Ala		
	290	295 300
Trp Asp Ala Gln Ile Arg Arg Ser Tyr Arg Pro Ser Val Val Arg Ala		

305 310 315 320
 Phe Gly Met Gly Trp Glu Lys Thr Val Ser His Ser Trp Thr Thr Leu
 325 330 335
 Lys Phe Phe Gly Lys Leu Ile Ser Gly Asn Ala Ser Val Ser His Ile
 340 345 350
 Ser Gly Pro Leu Thr Ile Ala Asp Ile Ala Gly Gln Ser Ala Glu Leu
 355 360 365
 Gly Leu Gln Ser Tyr Leu Glu Phe Leu Ala Leu Val Ser Ile Ser Leu
 370 375 380
 Gly Val Leu Asn Leu Leu Pro Val Pro Val Leu Asp Gly Gly His Leu
 385 390 395 400
 Val Phe Tyr Thr Ala Glu Trp Ile Arg Gly Lys Pro Leu Gly Glu Arg
 405 410 415
 Val Gln Asn Ile Gly Leu Arg Phe Gly Leu Ala Leu Met Met Leu Met
 420 425 430
 Met Ala Val Ala Phe Phe Asn Asp Val Thr Arg Leu Leu Gly
 435 440 445

<210> 185

<211> 1476

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1476)

<400> 185

atg aaa tac aaa gac ctg cgc gac ttc atc gcc atg ctc gag cag cag 48
 Met Lys Tyr Lys Asp Leu Arg Asp Phe Ile Ala Met Leu Glu Gln Gln

1

5

10

15

ggc aaa ctc aaa cgc atc gcg cac ccc gtt tcc ccg cat ttg gaa atg 96
 Gly Lys Leu Lys Arg Ile Ala His Pro Val Ser Pro His Leu Glu Met

20

25

30

acc gaa atc gcc gac cgc gtg ctg cgc gcc gaa ggg ccg gcg ttg ttg 144
 Thr Glu Ile Ala Asp Arg Val Leu Arg Ala Glu Gly Pro Ala Leu Leu

35	40	45	
ttt gaa cac cca gtt aag ccc gac ggt acg cgc tat gat tat ccc gtg			192
Phe Glu His Pro Val Lys Pro Asp Gly Thr Arg Tyr Asp Tyr Pro Val			
50	55	60	
ttg gca aac ctg ttc ggc acg ccc gaa cgt gtg gcg atg ggc atg ggc			240
Leu Ala Asn Leu Phe Gly Thr Pro Glu Arg Val Ala Met Gly Met Gly			
65	70	75	80
gcg gac agc gtg tcc aag ctg cgc gaa atc ggg cag acg ctg gcg tat			288
Ala Asp Ser Val Ser Lys Leu Arg Glu Ile Gly Gln Thr Leu Ala Tyr			
85	90	95	
ttg aaa gaa ccc gaa ccg ccc aaa ggc att aaa gac gcg ttt tcc aaa			336
Leu Lys Glu Pro Glu Pro Pro Lys Gly Ile Lys Asp Ala Phe Ser Lys			
100	105	110	
ctg ccg ctc ttg aaa gac att tgg agc atg gcg ccg aac gtg gtg aaa			384
Leu Pro Leu Leu Lys Asp Ile Trp Ser Met Ala Pro Asn Val Val Lys			
115	120	125	
aat gcg ccg tgt cag gaa atc gta tgg gaa ggc gaa gac gtt gat ttg			432
Asn Ala Pro Cys Gln Glu Ile Val Trp Glu Gly Glu Asp Val Asp Leu			
130	135	140	
tat caa ctt ccg att cag cat tgc tgg ccg gaa gac gtt gcg ccg ctg			480
Tyr Gln Leu Pro Ile Gln His Cys Trp Pro Glu Asp Val Ala Pro Leu			
145	150	155	160
gta acg tgg ggc ttg acc gtc acg cgc ggg ccg cac aaa aaa cgc caa			528
Val Thr Trp Gly Leu Thr Val Thr Arg Gly Pro His Lys Lys Arg Gln			
165	170	175	
aat ctc ggc att tac cgc caa caa tta atc ggc ata aac aag ctg att			576
Asn Leu Gly Ile Tyr Arg Gln Gln Leu Ile Gly Ile Asn Lys Leu Ile			
180	185	190	
atg cgc tgg ctg tcg cat cgc ggc ggc gcg ctg gat tat cag gaa ttc			624
Met Arg Trp Leu Ser His Arg Gly Gly Ala Leu Asp Tyr Gln Glu Phe			
195	200	205	
cgc aaa ctc aat ccc gat acg ccg tat ccc gtc gcc gtc gtg ctc ggc			672
Arg Lys Leu Asn Pro Asp Thr Pro Tyr Pro Val Ala Val Val Leu Gly			
210	215	220	
tgc gac cct gcc acc att ttg ggc gcg gta acg ccc gtt ccc gat acc			720
Cys Asp Pro Ala Thr Ile Leu Gly Ala Val Thr Pro Val Pro Asp Thr			

225	230	235	240	
ttg agc gaa tac cag ttt gcc gga ctg ctg cgc ggt tcg cgg acg gaa	768			
Leu Ser Glu Tyr Gln Phe Ala Gly Leu Leu Arg Gly Ser Arg Thr Glu				
245	250	255		
ctg gtg aaa tgt atc ggc aac gat ttg caa gtg cct gcc cgc gca gaa	816			
Leu Val Lys Cys Ile Gly Asn Asp Leu Gln Val Pro Ala Arg Ala Glu				
260	265	270		
atc gtg ttg gaa ggc gtc atc cat cca aac gaa acc gcg ttg gaa ggt	864			
Ile Val Leu Glu Gly Val Ile His Pro Asn Glu Thr Ala Leu Glu Gly				
275	280	285		
cca tac ggc gac cac acg ggc tat tac aac gag cag gac cat ttc ccc	912			
Pro Tyr Gly Asp His Thr Gly Tyr Tyr Asn Glu Gln Asp His Phe Pro				
290	295	300		
gtg ttt acg gtc gaa cgc atc acc atg cgc gaa aac ccg att tac cac	960			
Val Phe Thr Val Glu Arg Ile Thr Met Arg Glu Asn Pro Ile Tyr His				
305	310	315	320	
tct acc tac aca ggc aaa ccg ccc gat gaa ccc gcc gtt ttg ggc gtg	1008			
Ser Thr Tyr Thr Gly Lys Pro Pro Asp Glu Pro Ala Val Leu Gly Val				
325	330	335		
gcg ttg aac gaa gtg ttc gta ccg ctt ttg caa aag cag ttc ccc gaa	1056			
Ala Leu Asn Glu Val Phe Val Pro Leu Leu Gln Lys Gln Phe Pro Glu				
340	345	350		
atc acc gat ttt tac ctg ccg ccc gaa ggc tgc tcc tac cgc atg gcg	1104			
Ile Thr Asp Phe Tyr Leu Pro Pro Glu Gly Cys Ser Tyr Arg Met Ala				
355	360	365		
gtg gtg agc atg aaa aaa cag tac gcc gga cac gcc aag cgc gtg atg	1152			
Val Val Ser Met Lys Lys Gln Tyr Ala Gly His Ala Lys Arg Val Met				
370	375	380		
atg ggc tgc tgg tcg ttc ctg cgc cag ttt atg tac acc aaa ttc atc	1200			
Met Gly Cys Trp Ser Phe Leu Arg Gln Phe Met Tyr Thr Lys Phe Ile				
385	390	395	400	
att gtg gtg gat gac gat gtg gat gtg cgc gac tgg aaa gaa gtc atc	1248			
Ile Val Val Asp Asp Asp Val Asp Val Arg Asp Trp Lys Glu Val Ile				
405	410	415		
tgg gcg gta acc acg cgc atg gac ccc gtg cgc gat acc gtt ttg atg	1296			
Trp Ala Val Thr Thr Arg Met Asp Pro Val Arg Asp Thr Val Leu Met				

420	425	430	
gaa aac acg ccc atc gac tac ctc gac ttc gcc agc ccc gtc agc gga			1344
Glu Asn Thr Pro Ile Asp Tyr Leu Asp Phe Ala Ser Pro Val Ser Gly			
435	440	445	
ctt ggc ggc aaa atg ggt ttg gat gcg acc aac aag tgg ccg ggc gaa			1392
Leu Gly Gly Lys Met Gly Leu Asp Ala Thr Asn Lys Trp Pro Gly Glu			
450	455	460	
acc gac cgc gaa tgg gga cgg gtg att aaa aaa gac cct gcg gtt acg			1440
Thr Asp Arg Glu Trp Gly Arg Val Ile Lys Lys Asp Pro Ala Val Thr			
465	470	475	480
gct aag att gat gag att tgg gag gaa ttg ggg ttg			1476
Ala Lys Ile Asp Glu Ile Trp Glu Glu Leu Gly Leu			
485	490		

<210> 186

<211> 492

<212> PRT

<213> Neisseria meningitidis

<400> 186

Met Lys Tyr Lys Asp Leu Arg Asp Phe Ile Ala Met Leu Glu Gln Gln			
1	5	10	15
Gly Lys Leu Lys Arg Ile Ala His Pro Val Ser Pro His Leu Glu Met			
20	25	30	
Thr Glu Ile Ala Asp Arg Val Leu Arg Ala Glu Gly Pro Ala Leu Leu			
35	40	45	
Phe Glu His Pro Val Lys Pro Asp Gly Thr Arg Tyr Asp Tyr Pro Val			
50	55	60	
Leu Ala Asn Leu Phe Gly Thr Pro Glu Arg Val Ala Met Gly Met Gly			
65	70	75	80
Ala Asp Ser Val Ser Lys Leu Arg Glu Ile Gly Gln Thr Leu Ala Tyr			
85	90	95	
Leu Lys Glu Pro Glu Pro Pro Lys Gly Ile Lys Asp Ala Phe Ser Lys			
100	105	110	
Leu Pro Leu Leu Lys Asp Ile Trp Ser Met Ala Pro Asn Val Val Lys			
115	120	125	

Asn Ala Pro Cys Gln Glu Ile Val Trp Glu Gly Glu Asp Val Asp Leu
 130 135 140
 Tyr Gln Leu Pro Ile Gln His Cys Trp Pro Glu Asp Val Ala Pro Leu
 145 150 155 160
 Val Thr Trp Gly Leu Thr Val Thr Arg Gly Pro His Lys Lys Arg Gln
 165 170 175
 Asn Leu Gly Ile Tyr Arg Gln Gln Leu Ile Gly Ile Asn Lys Leu Ile
 180 185 190
 Met Arg Trp Leu Ser His Arg Gly Gly Ala Leu Asp Tyr Gln Glu Phe
 195 200 205
 Arg Lys Leu Asn Pro Asp Thr Pro Tyr Pro Val Ala Val Val Leu Gly
 210 215 220
 Cys Asp Pro Ala Thr Ile Leu Gly Ala Val Thr Pro Val Pro Asp Thr
 225 230 235 240
 Leu Ser Glu Tyr Gln Phe Ala Gly Leu Leu Arg Gly Ser Arg Thr Glu
 245 250 255
 Leu Val Lys Cys Ile Gly Asn Asp Leu Gln Val Pro Ala Arg Ala Glu
 260 265 270
 Ile Val Leu Glu Gly Val Ile His Pro Asn Glu Thr Ala Leu Glu Gly
 275 280 285
 Pro Tyr Gly Asp His Thr Gly Tyr Tyr Asn Glu Gln Asp His Phe Pro
 290 295 300
 Val Phe Thr Val Glu Arg Ile Thr Met Arg Glu Asn Pro Ile Tyr His
 305 310 315 320
 Ser Thr Tyr Thr Gly Lys Pro Pro Asp Glu Pro Ala Val Leu Gly Val
 325 330 335
 Ala Leu Asn Glu Val Phe Val Pro Leu Leu Gln Lys Gln Phe Pro Glu
 340 345 350
 Ile Thr Asp Phe Tyr Leu Pro Pro Glu Gly Cys Ser Tyr Arg Met Ala
 355 360 365
 Val Val Ser Met Lys Lys Gln Tyr Ala Gly His Ala Lys Arg Val Met
 370 375 380

Met Gly Cys Trp Ser Phe Leu Arg Gln Phe Met Tyr Thr Lys Phe Ile
 385 390 395 400

Ile Val Val Asp Asp Asp Val Asp Val Arg Asp Trp Lys Glu Val Ile
 405 410 415

Trp Ala Val Thr Thr Arg Met Asp Pro Val Arg Asp Thr Val Leu Met
 420 425 430

Glu Asn Thr Pro Ile Asp Tyr Leu Asp Phe Ala Ser Pro Val Ser Gly
 435 440 445

Leu Gly Gly Lys Met Gly Leu Asp Ala Thr Asn Lys Trp Pro Gly Glu
 450 455 460

Thr Asp Arg Glu Trp Gly Arg Val Ile Lys Lys Asp Pro Ala Val Thr
 465 470 475 480

Ala Lys Ile Asp Glu Ile Trp Glu Glu Leu Gly Leu
 485 490

<210> 187

<211> 1191

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1191)

<400> 187

atg ttc ttc aag cac atc gaa gcc gcc ccc gcc gat ccg att ctc ggt 48
 Met Phe Phe Lys His Ile Glu Ala Ala Pro Ala Asp Pro Ile Leu Gly
 1 5 10 15

ttg ggc gaa gca ttc aaa gcc gaa acc cgc ccc gaa aaa gtc aac ctc 96
 Leu Gly Glu Ala Phe Lys Ala Glu Thr Arg Pro Glu Lys Val Asn Leu
 20 25 30

ggc atc ggc gtg tac aaa gac gca tcc ggc gcg aca cct att gtc aaa 144
 Gly Ile Gly Val Tyr Lys Asp Ala Ser Gly Ala Thr Pro Ile Val Lys
 35 40 45

gcc gtc aaa gaa gcc gaa aaa cgc ctg ttg gaa agc gaa aca act aaa 192
 Ala Val Lys Glu Ala Glu Lys Arg Leu Leu Glu Ser Glu Thr Thr Lys

50	55	60	
aac tac ctc acc atc gac ggc gtt gcc gac tac aac gag caa acc caa			240
Asn Tyr Leu Thr Ile Asp Gly Val Ala Asp Tyr Asn Glu Gln Thr Gln			
65	70	75	80
att ctg ctg ttc ggc aaa gac cac gaa atc atc gcc agc cgt cgc gcc			288
Ile Leu Leu Phe Gly Lys Asp His Glu Ile Ile Ala Ser Arg Arg Ala			
	85	90	95
aaa aca gcg caa agc ctt ggc ggt aca ggc gca ttg cgt att gcg gcc			336
Lys Thr Ala Gln Ser Leu Gly Gly Thr Gly Ala Leu Arg Ile Ala Ala			
	100	105	110
gag ttt gcc aaa cgt cag ttg aac gcg caa acc atc tgg att tcc aat			384
Glu Phe Ala Lys Arg Gln Leu Asn Ala Gln Thr Ile Trp Ile Ser Asn			
	115	120	125
ccg act tgg ccc aac cac aac gcc atc gcc aaa gcg gtc ggt atc caa			432
Pro Thr Trp Pro Asn His Asn Ala Ile Ala Lys Ala Val Gly Ile Gln			
	130	135	140
gac caa cct tat cgc tac tat gat gcc gcc aaa cac ggt ttg gat tgg			480
Asp Gln Pro Tyr Arg Tyr Tyr Asp Ala Ala Lys His Gly Leu Asp Trp			
145	150	155	160
gac ggt atg att gaa gac ttg agc caa gcg caa aaa ggc gac atc gtc			528
Asp Gly Met Ile Glu Asp Leu Ser Gln Ala Gln Lys Gly Asp Ile Val			
	165	170	175
ctg ctg cac ggc tgc tgc cac aac cct acc ggt atc gac cct acg ccc			576
Leu Leu His Gly Cys Cys His Asn Pro Thr Gly Ile Asp Pro Thr Pro			
	180	185	190
gaa caa tgg gaa act ctg gca aaa ctt tct gcc gaa aaa ggc tgg tta			624
Glu Gln Trp Glu Thr Leu Ala Lys Leu Ser Ala Glu Lys Gly Trp Leu			
	195	200	205
ccg ctg ttt gac ttt gcc tac caa ggc ttc ggc aat ggt ttg gaa gaa			672
Pro Leu Phe Asp Phe Ala Tyr Gln Gly Phe Gly Asn Gly Leu Glu Glu			
210	215	220	
gat gcc tac ggc ctg cgc gtg ttc ttg aaa cac aat aca gaa ttg ctg			720
Asp Ala Tyr Gly Leu Arg Val Phe Leu Lys His Asn Thr Glu Leu Leu			
225	230	235	240
att gcc agc tct tat tcc aaa aac ttc ggt atg tac aac gag cgc gtc			768
Ile Ala Ser Ser Tyr Ser Lys Asn Phe Gly Met Tyr Asn Glu Arg Val			

245	250	255	
ggc gcg ttc act ttg gtg gcc gaa gat gaa gca acc gcc gcc cgc gcc			816
Gly Ala Phe Thr Leu Val Ala Glu Asp Glu Ala Thr Ala Ala Arg Ala			
260	265	270	
cac agc caa gtc aaa acc att atc cgt acc ttg tat tcc aac cct gct			864
His Ser Gln Val Lys Thr Ile Ile Arg Thr Leu Tyr Ser Asn Pro Ala			
275	280	285	
tca cac ggc gcg aac act att gcg ctg gtg ttg aaa aat gat gat ttg			912
Ser His Gly Ala Asn Thr Ile Ala Leu Val Leu Lys Asn Asp Asp Leu			
290	295	300	
aaa gca caa tgg att gcc gaa ctc gat gaa atg cgc gcc cgt atc aaa			960
Lys Ala Gln Trp Ile Ala Glu Leu Asp Glu Met Arg Gly Arg Ile Lys			
305	310	315	320
gcc atg cgc caa aaa ttt gtc gag ttg ctc aaa gcc aaa ggt gca act			1008
Ala Met Arg Gln Lys Phe Val Glu Leu Leu Lys Ala Lys Gly Ala Thr			
325	330	335	
caa gac ttt gat ttc att atc gaa caa aac gcc atg ttc tcc ttc agc			1056
Gln Asp Phe Asp Phe Ile Ile Glu Gln Asn Gly Met Phe Ser Phe Ser			
340	345	350	
ggc ttg act ccc gaa caa gtc gac cgt tta aaa aac gag ttt gcc att			1104
Gly Leu Thr Pro Glu Gln Val Asp Arg Leu Lys Asn Glu Phe Ala Ile			
355	360	365	
tat gcc gtc cgc tcc gcc cgc atc aac gtc gcc gcc att acc gac gac			1152
Tyr Ala Val Arg Ser Gly Arg Ile Asn Val Ala Gly Ile Thr Asp Asp			
370	375	380	
aac atc gat tat ctg tgc gaa agt atc gta aaa gta ctg			1191
Asn Ile Asp Tyr Leu Cys Glu Ser Ile Val Lys Val Leu			
385	390	395	

<210> 188

<211> 397

<212> PRT

<213> Neisseria meningitidis

<400> 188

Met Phe Phe Lys His Ile Glu Ala Ala Pro Ala Asp Pro Ile Leu Gly

1

5

10

15

Leu Gly Glu Ala Phe Lys Ala Glu Thr Arg Pro Glu Lys Val Asn Leu
 20 25 30

Gly Ile Gly Val Tyr Lys Asp Ala Ser Gly Ala Thr Pro Ile Val Lys
 35 40 45

Ala Val Lys Glu Ala Glu Lys Arg Leu Leu Glu Ser Glu Thr Thr Lys
 50 55 60

Asn Tyr Leu Thr Ile Asp Gly Val Ala Asp Tyr Asn Glu Gln Thr Gln
 65 70 75 80

Ile Leu Leu Phe Gly Lys Asp His Glu Ile Ile Ala Ser Arg Arg Ala
 85 90 95

Lys Thr Ala Gln Ser Leu Gly Gly Thr Gly Ala Leu Arg Ile Ala Ala
 100 105 110

Glu Phe Ala Lys Arg Gln Leu Asn Ala Gln Thr Ile Trp Ile Ser Asn
 115 120 125

Pro Thr Trp Pro Asn His Asn Ala Ile Ala Lys Ala Val Gly Ile Gln
 130 135 140

Asp Gln Pro Tyr Arg Tyr Tyr Asp Ala Ala Lys His Gly Leu Asp Trp
 145 150 155 160

Asp Gly Met Ile Glu Asp Leu Ser Gln Ala Gln Lys Gly Asp Ile Val
 165 170 175

Leu Leu His Gly Cys Cys His Asn Pro Thr Gly Ile Asp Pro Thr Pro
 180 185 190

Glu Gln Trp Glu Thr Leu Ala Lys Leu Ser Ala Glu Lys Gly Trp Leu
 195 200 205

Pro Leu Phe Asp Phe Ala Tyr Gln Gly Phe Gly Asn Gly Leu Glu Glu
 210 215 220

Asp Ala Tyr Gly Leu Arg Val Phe Leu Lys His Asn Thr Glu Leu Leu
 225 230 235 240

Ile Ala Ser Ser Tyr Ser Lys Asn Phe Gly Met Tyr Asn Glu Arg Val
 245 250 255

Gly Ala Phe Thr Leu Val Ala Glu Asp Glu Ala Thr Ala Ala Arg Ala
 260 265 270

His Ser Gln Val Lys Thr Ile Ile Arg Thr Leu Tyr Ser Asn Pro Ala
 275 280 285
 Ser His Gly Ala Asn Thr Ile Ala Leu Val Leu Lys Asn Asp Asp Leu
 290 295 300
 Lys Ala Gln Trp Ile Ala Glu Leu Asp Glu Met Arg Gly Arg Ile Lys
 305 310 315 320
 Ala Met Arg Gln Lys Phe Val Glu Leu Leu Lys Ala Lys Gly Ala Thr
 325 330 335
 Gln Asp Phe Asp Phe Ile Ile Glu Gln Asn Gly Met Phe Ser Phe Ser
 340 345 350
 Gly Leu Thr Pro Glu Gln Val Asp Arg Leu Lys Asn Glu Phe Ala Ile
 355 360 365
 Tyr Ala Val Arg Ser Gly Arg Ile Asn Val Ala Gly Ile Thr Asp Asp
 370 375 380
 Asn Ile Asp Tyr Leu Cys Glu Ser Ile Val Lys Val Leu
 385 390 395

<210> 189
 <211> 462
 <212> DNA
 <213> Neisseria meningitidis

<220>
 <221> CDS
 <222> (1)..(462)

<400> 189
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 Leu Leu Cys Pro Glu Lys Met Ser Gly Met Ala Gly Gln Tyr Pro Tyr
 1 5 10 15
 ggc gtc cgg tcg ggt ttg cgg agg aac ggc ttg aaa ctt tgg gat att 96
 Gly Val Arg Ser Gly Leu Arg Arg Asn Gly Leu Lys Leu Trp Asp Ile
 20 25 30
 cat ttt aga atg acc cgt ttt atc gtc gca aga tgc ggt tta ttg ttt 144
 His Phe Arg Met Thr Arg Phe Ile Val Ala Arg Cys Gly Leu Leu Phe
 35 40 45

gca acc ctt aaa gga aaa acc atg aag aaa atg ttc gtg ctg ttc tgt 192
 Ala Thr Leu Lys Gly Lys Thr Met Lys Lys Met Phe Val Leu Phe Cys
 50 55 60

atg ctg ttc tcc tgc gcc ttc tcc ctt gcg gcg gta aac atc aat gcg 240
 Met Leu Phe Ser Cys Ala Phe Ser Leu Ala Ala Val Asn Ile Asn Ala
 65 70 75 80

gct tcg cag cag gag ctg gag gcg ctg ccg ggc ata ggc cct gcg aag 288
 Ala Ser Gln Gln Glu Leu Glu Ala Leu Pro Gly Ile Gly Pro Ala Lys
 85 90 95

gcg aag gcc att gcg gaa tac cgt gcg caa aac ggt gcg ttc aag tct 336
 Ala Lys Ala Ile Ala Glu Tyr Arg Ala Gln Asn Gly Ala Phe Lys Ser
 100 105 110

gta gac gat ttg acc aag gta aag ggc atc ggc cct gcg gtg ctg gcg 384
 Val Asp Asp Leu Thr Lys Val Lys Gly Ile Gly Pro Ala Val Leu Ala
 115 120 125

aag ctg aag gat cag gct tct gtc ggt gcg ccc gca cca aaa ggc ccg 432
 Lys Leu Lys Asp Gln Ala Ser Val Gly Ala Pro Ala Pro Lys Gly Pro
 130 135 140

gca aaa ccg gct ctg ccc gca gct aaa aaa 462
 Ala Lys Pro Ala Leu Pro Ala Ala Lys Lys
 145 150

<210> 190

<211> 154

<212> PRT

<213> Neisseria meningitidis

<400> 190

Leu Leu Cys Pro Glu Lys Met Ser Gly Met Ala Gly Gln Tyr Pro Tyr
 1 5 10 15

Gly Val Arg Ser Gly Leu Arg Arg Asn Gly Leu Lys Leu Trp Asp Ile
 20 25 30

His Phe Arg Met Thr Arg Phe Ile Val Ala Arg Cys Gly Leu Leu Phe
 35 40 45

Ala Thr Leu Lys Gly Lys Thr Met Lys Lys Met Phe Val Leu Phe Cys
 50 55 60

Met Leu Phe Ser Cys Ala Phe Ser Leu Ala Ala Val Asn Ile Asn Ala

<400> 191																
atg	ttt	gct	ttt	tta	gaa	gcc	ttt	ttt	gtc	gaa	tac	ggg	tat	gcg	gct	48
Met	Phe	Ala	Phe	Leu	Glu	Ala	Phe	Phe	Val	Glu	Tyr	Gly	Tyr	Ala	Ala	
1		5			10					15						
gtt	ttt	ttt	gta	ttg	gtc	atc	tgc	ggg	ttc	ggc	gtg	ccg	att	ccc	gag	96
Val	Phe	Phe	Val	Leu	Val	Ile	Cys	Gly	Phe	Gly	Val	Pro	Ile	Pro	Glu	
			20		25					30						
gat	ttg	acc	ttg	gta	aca	ggc	ggc	gtg	att	tcg	ggg	atg	ggg	tat	acc	144
Asp	Leu	Thr	Leu	Val	Thr	Gly	Gly	Val	Ile	Ser	Gly	Met	Gly	Tyr	Thr	
35			40					45								
aat	ccg	cat	att	atg	ttt	gca	gtc	ggg	atg	ctc	ggc	gta	ttg	gtc	ggg	192
Asn	Pro	His	Ile	Met	Phe	Ala	Val	Gly	Met	Leu	Gly	Val	Leu	Val	Gly	
50			55					60								
gac	ggc	atc	atg	ttc	gcc	gcc	gga	cga	att	tgg	ggg	cag	aaa	atc	cta	240
Asp	Gly	Ile	Met	Phe	Ala	Ala	Gly	Arg	Ile	Trp	Gly	Gln	Lys	Ile	Leu	
65		70			75					80						

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agg ttc aaa cct att gcg cgc atc atg acg ccg aaa cgt tat gag cag 288
Arg Phe Lys Pro Ile Ala Arg Ile Met Thr Pro Lys Arg Tyr Glu Gln
      85                      90                      95

ggt cag gaa aaa ttc gac aaa tac ggt aac tgg gtc tta ttt gtc gcc 336
Val Gln Glu Lys Phe Asp Lys Tyr Gly Asn Trp Val Leu Phe Val Ala
      100                     105                     110

cgt ttc ctg ccc ggt ttg aga acg gcc gta ttt gtt aca gcc ggt atc 384
Arg Phe Leu Pro Gly Leu Arg Thr Ala Val Phe Val Thr Ala Gly Ile
      115                     120                     125

agc cgc aag gtt tca tac ttg cgt ttt atc att atg gat gga ctg gcc 432
Ser Arg Lys Val Ser Tyr Leu Arg Phe Ile Ile Met Asp Gly Leu Ala
      130                     135                     140

gca ctg att tcc gtc cct att tgg att tat ctg ggc gaa tac ggt gcg 480
Ala Leu Ile Ser Val Pro Ile Trp Ile Tyr Leu Gly Glu Tyr Gly Ala
      145                     150                     155                     160

cac aac atc gat tgg ctg atg gcg aaa atg cac agc ctg caa tcg ggt 528
His Asn Ile Asp Trp Leu Met Ala Lys Met His Ser Leu Gln Ser Gly
      165                     170                     175

att ttt gtt atc ttg ggt ata ggt gcg acc gtt gtc gct tgg att tgg 576
Ile Phe Val Ile Leu Gly Ile Gly Ala Thr Val Val Ala Trp Ile Trp
      180                     185                     190

tgg aaa aaa cgc caa cgt atc cag ttt tac cgc agc aaa ttg aaa gaa 624
Trp Lys Lys Arg Gln Arg Ile Gln Phe Tyr Arg Ser Lys Leu Lys Glu
      195                     200                     205

aag cgg gcg caa cgc aaa gcc gcc aag gca gcc aaa aaa gcc gcg caa 672
Lys Arg Ala Gln Arg Lys Ala Ala Lys Ala Ala Lys Lys Ala Ala Gln
      210                     215                     220

agc aaa caa taa 684
Ser Lys Gln
225

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<210> 192

<211> 227

<212> PRT

<213> Neisseria meningitidis

<400> 192

Met Phe Ala Phe Leu Glu Ala Phe Phe Val Glu Tyr Gly Tyr Ala Ala
 1 5 10 15
 Val Phe Phe Val Leu Val Ile Cys Gly Phe Gly Val Pro Ile Pro Glu
 20 25 30
 Asp Leu Thr Leu Val Thr Gly Gly Val Ile Ser Gly Met Gly Tyr Thr
 35 40 45
 Asn Pro His Ile Met Phe Ala Val Gly Met Leu Gly Val Leu Val Gly
 50 55 60
 Asp Gly Ile Met Phe Ala Ala Gly Arg Ile Trp Gly Gln Lys Ile Leu
 65 70 75 80
 Arg Phe Lys Pro Ile Ala Arg Ile Met Thr Pro Lys Arg Tyr Glu Gln
 85 90 95
 Val Gln Glu Lys Phe Asp Lys Tyr Gly Asn Trp Val Leu Phe Val Ala
 100 105 110
 Arg Phe Leu Pro Gly Leu Arg Thr Ala Val Phe Val Thr Ala Gly Ile
 115 120 125
 Ser Arg Lys Val Ser Tyr Leu Arg Phe Ile Ile Met Asp Gly Leu Ala
 130 135 140
 Ala Leu Ile Ser Val Pro Ile Trp Ile Tyr Leu Gly Glu Tyr Gly Ala
 145 150 155 160
 His Asn Ile Asp Trp Leu Met Ala Lys Met His Ser Leu Gln Ser Gly
 165 170 175
 Ile Phe Val Ile Leu Gly Ile Gly Ala Thr Val Val Ala Trp Ile Trp
 180 185 190
 Trp Lys Lys Arg Gln Arg Ile Gln Phe Tyr Arg Ser Lys Leu Lys Glu
 195 200 205
 Lys Arg Ala Gln Arg Lys Ala Ala Lys Ala Ala Lys Lys Ala Ala Gln
 210 215 220
 Ser Lys Gln
 225

<210> 193

<211> 1089

<212> DNA

<213> *Neisseria meningitidis*

<220>

<221> CDS

<222> (1)..(1089)

<400> 193

atg aat gac tac acg cag cag ctt caa ggt aaa aaa gac tac ctt aaa 48
 Met Asn Asp Tyr Thr Gln Gln Leu Gln Gly Lys Lys Asp Tyr Leu Lys
 1 5 10 15

acc ctt ttt gca ggt ttg gat gtt cct gag tgg gaa gtg tac gaa tct 96
 Thr Leu Phe Ala Gly Leu Asp Val Pro Glu Trp Glu Val Tyr Glu Ser
 20 25 30

ccg gac aaa cat tac cgt atg cgt gcc gag ttc cgt att tgg cac gaa 144
 Pro Asp Lys His Tyr Arg Met Arg Ala Glu Phe Arg Ile Trp His Glu
 35 40 45

ggc ggg gaa atg ttt tat gcc atg ttt gaa aaa ggg cag aaa gcc agc 192
 Gly Gly Glu Met Phe Tyr Ala Met Phe Glu Lys Gly Gln Lys Ala Ser
 50 55 60

ggc gca agc atg ata cgc tgc gac cgt ttt gaa gca gct tcc gag gct 240
 Gly Ala Ser Met Ile Arg Cys Asp Arg Phe Glu Ala Ala Ser Glu Ala
 65 70 75 80

gtc aac cgc ctc atg ccc gag ctg atc gcc gcc gcc gcg caa tcc ccc 288
 Val Asn Arg Leu Met Pro Glu Leu Ile Ala Ala Ala Ala Gln Ser Pro
 85 90 95

gaa ctc aaa aaa cgc tgg tat gcc gtc gaa ttt ctg tcc acg ctc agc 336
 Glu Leu Lys Lys Arg Trp Tyr Ala Val Glu Phe Leu Ser Thr Leu Ser
 100 105 110

ggg gaa atg ctg gtt acc atg att tac cac aaa agg ctt gat gct gag 384
 Gly Glu Met Leu Val Thr Met Ile Tyr His Lys Arg Leu Asp Ala Glu
 115 120 125

tgg atg cag gcg gcg caa gcg tta cag caa cag ttg gat att tcc gtt 432
 Trp Met Gln Ala Ala Gln Ala Leu Gln Gln Gln Leu Asp Ile Ser Val
 130 135 140

att ggg cgg agc agg gga cag aaa ata gtc tta aaa cag gac tat gta 480
 Ile Gly Arg Ser Arg Gly Gln Lys Ile Val Leu Lys Gln Asp Tyr Val
 145 150 155 160


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acg gaa act ttg aag gtc ggc aac cgg gat ttc cgt tat cgg caa att 528
Thr Glu Thr Leu Lys Val Gly Asn Arg Asp Phe Arg Tyr Arg Gln Ile
      165              170              175

gaa ggc agt ttt acc caa ccg aat gcc gcc gtg tgt cag aaa atg ctt 576
Glu Gly Ser Phe Thr Gln Pro Asn Ala Ala Val Cys Gln Lys Met Leu
      180              185              190

gaa tgg gcg tgc cgc acg gcg gaa ggc ttg ggc agc gac ctg ctc gaa 624
Glu Trp Ala Cys Arg Thr Ala Glu Gly Leu Gly Ser Asp Leu Leu Glu
      195              200              205

ctg tat tgc gga aac ggc aat ttc acc ctg ccg ctg tcc cga tat ttc 672
Leu Tyr Cys Gly Asn Gly Asn Phe Thr Leu Pro Leu Ser Arg Tyr Phe
      210              215              220

cga cag gtt ttg gca acc gaa atc tcc aaa acc tcc gtc agc gcg gcg 720
Arg Gln Val Leu Ala Thr Glu Ile Ser Lys Thr Ser Val Ser Ala Ala
      225              230              235              240

caa tgg aat atc gaa gcc aac cgc atc ggc aac atc aaa atc gcc cgt 768
Gln Trp Asn Ile Glu Ala Asn Arg Ile Gly Asn Ile Lys Ile Ala Arg
      245              250              255

ctg tct gct gaa gaa ttt acc gaa gcc tat acc gga aaa cgt gag ttt 816
Leu Ser Ala Glu Glu Phe Thr Glu Ala Tyr Thr Gly Lys Arg Glu Phe
      260              265              270

aaa cgc ctt aag gat ggc ggt att gct ttg acg gat tac gca ttt tcc 864
Lys Arg Leu Lys Asp Gly Gly Ile Ala Leu Thr Asp Tyr Ala Phe Ser
      275              280              285

act att ttc gtc gat ccg ccg cgt gcc ggt atc gat gag gaa act ttg 912
Thr Ile Phe Val Asp Pro Pro Arg Ala Gly Ile Asp Glu Glu Thr Leu
      290              295              300

aag ctg gtg tcg cag ttt gac aac atc atc tac att tcc tgc aat ccc 960
Lys Leu Val Ser Gln Phe Asp Asn Ile Ile Tyr Ile Ser Cys Asn Pro
      305              310              315              320

gaa acg ttg cgc gcc aat ttg gac acc ttg gcg gaa acc cat gcg gtc 1008
Glu Thr Leu Arg Ala Asn Leu Asp Thr Leu Ala Glu Thr His Ala Val
      325              330              335

gaa cgt gcc gcc ctg ttc gac caa ttc ccg ttt acg cac cat atc gaa 1056
Glu Arg Ala Ala Leu Phe Asp Gln Phe Pro Phe Thr His His Ile Glu
      340              345              350

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agt ggt gta ttg ttg aaa aag aaa atc ctt tga 1089
 Ser Gly Val Leu Leu Lys Lys Lys Ile Leu
 355 360

<210> 194
 <211> 362
 <212> PRT
 <213> Neisseria meningitidis

<400> 194
 Met Asn Asp Tyr Thr Gln Gln Leu Gln Gly Lys Lys Asp Tyr Leu Lys
 1 5 10 15
 Thr Leu Phe Ala Gly Leu Asp Val Pro Glu Trp Glu Val Tyr Glu Ser
 20 25 30
 Pro Asp Lys His Tyr Arg Met Arg Ala Glu Phe Arg Ile Trp His Glu
 35 40 45
 Gly Gly Glu Met Phe Tyr Ala Met Phe Glu Lys Gly Gln Lys Ala Ser
 50 55 60
 Gly Ala Ser Met Ile Arg Cys Asp Arg Phe Glu Ala Ala Ser Glu Ala
 65 70 75 80
 Val Asn Arg Leu Met Pro Glu Leu Ile Ala Ala Ala Ala Gln Ser Pro
 85 90 95
 Glu Leu Lys Lys Arg Trp Tyr Ala Val Glu Phe Leu Ser Thr Leu Ser
 100 105 110
 Gly Glu Met Leu Val Thr Met Ile Tyr His Lys Arg Leu Asp Ala Glu
 115 120 125
 Trp Met Gln Ala Ala Gln Ala Leu Gln Gln Gln Leu Asp Ile Ser Val
 130 135 140
 Ile Gly Arg Ser Arg Gly Gln Lys Ile Val Leu Lys Gln Asp Tyr Val
 145 150 155 160
 Thr Glu Thr Leu Lys Val Gly Asn Arg Asp Phe Arg Tyr Arg Gln Ile
 165 170 175
 Glu Gly Ser Phe Thr Gln Pro Asn Ala Ala Val Cys Gln Lys Met Leu
 180 185 190

Glu Trp Ala Cys Arg Thr Ala Glu Gly Leu Gly Ser Asp Leu Leu Glu
 195 200 205

Leu Tyr Cys Gly Asn Gly Asn Phe Thr Leu Pro Leu Ser Arg Tyr Phe
 210 215 220

Arg Gln Val Leu Ala Thr Glu Ile Ser Lys Thr Ser Val Ser Ala Ala
 225 230 235 240

Gln Trp Asn Ile Glu Ala Asn Arg Ile Gly Asn Ile Lys Ile Ala Arg
 245 250 255

Leu Ser Ala Glu Glu Phe Thr Glu Ala Tyr Thr Gly Lys Arg Glu Phe
 260 265 270

Lys Arg Leu Lys Asp Gly Gly Ile Ala Leu Thr Asp Tyr Ala Phe Ser
 275 280 285

Thr Ile Phe Val Asp Pro Pro Arg Ala Gly Ile Asp Glu Glu Thr Leu
 290 295 300

Lys Leu Val Ser Gln Phe Asp Asn Ile Ile Tyr Ile Ser Cys Asn Pro
 305 310 315 320

Glu Thr Leu Arg Ala Asn Leu Asp Thr Leu Ala Glu Thr His Ala Val
 325 330 335

Glu Arg Ala Ala Leu Phe Asp Gln Phe Pro Phe Thr His His Ile Glu
 340 345 350

Ser Gly Val Leu Leu Lys Lys Lys Ile Leu
 355 360

<210> 195

<211> 2145

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(2145)

<400> 195

atg aat acc cca ttg ttc cgt ctc agc ctg ctc tcg ctt acc ctg gcg 48
 Met Asn Thr Pro Leu Phe Arg Leu Ser Leu Leu Ser Leu Thr Leu Ala
 1 5 10 15

gca ggt ttt gcc cat gcg gca gaa aat aat gcc aag gtc gta ctg gat 96
 Ala Gly Phe Ala His Ala Ala Glu Asn Asn Ala Lys Val Val Leu Asp
 20 25 30

acc gtt acc gta aaa ggc gac cgc caa ggc agc aaa atc cgt acc aac 144
 Thr Val Thr Val Lys Gly Asp Arg Gln Gly Ser Lys Ile Arg Thr Asn
 35 40 45

atc gtt acg ctg caa caa aaa gac gaa agc acc gca acc gat atg cgc 192
 Ile Val Thr Leu Gln Gln Lys Asp Glu Ser Thr Ala Thr Asp Met Arg
 50 55 60

gaa ctc tta aaa gaa gag ccc tcc atc gat ttc ggc ggc ggc aac ggc 240
 Glu Leu Leu Lys Glu Glu Pro Ser Ile Asp Phe Gly Gly Gly Asn Gly
 65 70 75 80

acg tcc caa ttc ctg acg ctg cgc ggc atg ggt caa aac tct gtc gac 288
 Thr Ser Gln Phe Leu Thr Leu Arg Gly Met Gly Gln Asn Ser Val Asp
 85 90 95

atc aag gtg gac aac gcc tat tcc gac agc caa atc ctt tac cac caa 336
 Ile Lys Val Asp Asn Ala Tyr Ser Asp Ser Gln Ile Leu Tyr His Gln
 100 105 110

ggc aga ttt att gtc gat ccc gct ttg gtt aaa gtc gtt tcc gta caa 384
 Gly Arg Phe Ile Val Asp Pro Ala Leu Val Lys Val Val Ser Val Gln
 115 120 125

aaa ggc gcg ggt tcc gcc tct gcc ggt atc ggc gcg acc aac ggc gcg 432
 Lys Gly Ala Gly Ser Ala Ser Ala Gly Ile Gly Ala Thr Asn Gly Ala
 130 135 140

atc atc acc aaa acc gtc gat gcc caa gac ctg ctc aaa ggc ttg gat 480
 Ile Ile Thr Lys Thr Val Asp Ala Gln Asp Leu Leu Lys Gly Leu Asp
 145 150 155 160

aaa aac tgg ggc gtg cgc ctc aac agc ggc ttt gcc agc aac gaa ggc 528
 Lys Asn Trp Gly Val Arg Leu Asn Ser Gly Phe Ala Ser Asn Glu Gly
 165 170 175

gta agc tac ggc gca agc gta ttc ggg aaa gag ggc aac ttc gac ggc 576
 Val Ser Tyr Gly Ala Ser Val Phe Gly Lys Glu Gly Asn Phe Asp Gly
 180 185 190

ttg ttc tct tac aac cgc aac aat gaa aaa gat tac gaa gca ggt aaa 624
 Leu Phe Ser Tyr Asn Arg Asn Asn Glu Lys Asp Tyr Glu Ala Gly Lys
 195 200 205

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ggc ttc cgt aat aat ttc aac ggc ggc aaa acc gta ccg tac agc gcg 672
Gly Phe Arg Asn Asn Phe Asn Gly Gly Lys Thr Val Pro Tyr Ser Ala
210 215 220

ctg gac aaa cgc agc tac ctc gcc aaa atc gga aca agc ttc ggc gac 720
Leu Asp Lys Arg Ser Tyr Leu Ala Lys Ile Gly Thr Ser Phe Gly Asp
225 230 235 240

ggc gac cac cgc atc gta ttg agc cat atg aaa gac cag cac cgg ggc 768
Gly Asp His Arg Ile Val Leu Ser His Met Lys Asp Gln His Arg Gly
245 250 255

atc cgt acc gtc cgt gaa gaa ttt acc gtc ggc ggc gat aaa gag cga 816
Ile Arg Thr Val Arg Glu Glu Phe Thr Val Gly Gly Asp Lys Glu Arg
260 265 270

ata agt atg gaa cgc caa gcc cct gct tac cgc gaa acc aca caa tcc 864
Ile Ser Met Glu Arg Gln Ala Pro Ala Tyr Arg Glu Thr Thr Gln Ser
275 280 285

aac acc aat ttg gcg tac acg ggt aaa aac ctg ggc ttt gtc gaa aaa 912
Asn Thr Asn Leu Ala Tyr Thr Gly Lys Asn Leu Gly Phe Val Glu Lys
290 295 300

ctg gat gcc aac gcc tat gtg ttg gaa aaa gaa cgc tat tcc gcc gat 960
Leu Asp Ala Asn Ala Tyr Val Leu Glu Lys Glu Arg Tyr Ser Ala Asp
305 310 315 320

gac agc ggc acc ggt tac gca ggc aat gta aaa ggc ccc aac cat acc 1008
Asp Ser Gly Thr Gly Tyr Ala Gly Asn Val Lys Gly Pro Asn His Thr
325 330 335

caa atc acc act cgg ggt atg aac ttc aac ttc gac agc cgc ctt gcc 1056
Gln Ile Thr Thr Arg Gly Met Asn Phe Asn Phe Asp Ser Arg Leu Ala
340 345 350

gaa caa acc ctg ctg aaa tac ggt atc aac tac cgc cat cag gaa atc 1104
Glu Gln Thr Leu Leu Lys Tyr Gly Ile Asn Tyr Arg His Gln Glu Ile
355 360 365

aaa ccg caa gcg ttt ttg aat tca caa ttt aaa att gaa gat aaa gaa 1152
Lys Pro Gln Ala Phe Leu Asn Ser Gln Phe Lys Ile Glu Asp Lys Glu
370 375 380

aaa gca act gat gaa gag aaa aat aag aac cgt gaa aat gaa aaa att 1200
Lys Ala Thr Asp Glu Glu Lys Asn Lys Asn Arg Glu Asn Glu Lys Ile
385 390 395 400

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gcc aaa gcc tac cgt ctg acc aac ccg acc aaa acc gat acc ggc gcg	1248
Ala Lys Ala Tyr Arg Leu Thr Asn Pro Thr Lys Thr Asp Thr Gly Ala	
405 410 415	
tat atc gaa gcc att cac gag att gac ggc ttt acc ctg acc ggc ggg	1296
Tyr Ile Glu Ala Ile His Glu Ile Asp Gly Phe Thr Leu Thr Gly Gly	
420 425 430	
ctg cgt tac gac cgc ttc aag gtg aaa acc cac gac ggc aaa acc gtt	1344
Leu Arg Tyr Asp Arg Phe Lys Val Lys Thr His Asp Gly Lys Thr Val	
435 440 445	
tca agc aac aac ctt aac ccg agt ttc ggc gtg att tgg cag ccg cac	1392
Ser Ser Asn Asn Leu Asn Pro Ser Phe Gly Val Ile Trp Gln Pro His	
450 455 460	
gaa cac tgg agc ttc agc gcg agc cac aac tac gcc agc cgc agc ccg	1440
Glu His Trp Ser Phe Ser Ala Ser His Asn Tyr Ala Ser Arg Ser Pro	
465 470 475 480	
cgc ctg tat gac gcg ctg caa acc cac ggc aaa cgc ggc atc atc tcg	1488
Arg Leu Tyr Asp Ala Leu Gln Thr His Gly Lys Arg Gly Ile Ile Ser	
485 490 495	
att gcc gac ggc acg aaa gcc gaa cgc gcg cgc aat acc gaa atc ggc	1536
Ile Ala Asp Gly Thr Lys Ala Glu Arg Ala Arg Asn Thr Glu Ile Gly	
500 505 510	
ttc aac tac aac gac ggc acg ttt gcc gca aac ggc agc tac ttc tgg	1584
Phe Asn Tyr Asn Asp Gly Thr Phe Ala Ala Asn Gly Ser Tyr Phe Trp	
515 520 525	
cag acc atc aaa gac gcg ctt gcc aat ccg caa aac cgc cac gac tct	1632
Gln Thr Ile Lys Asp Ala Leu Ala Asn Pro Gln Asn Arg His Asp Ser	
530 535 540	
gtc gcc gtc cgt gaa gcc gtc aat gcc ggt tac atc aaa aac cac ggt	1680
Val Ala Val Arg Glu Ala Val Asn Ala Gly Tyr Ile Lys Asn His Gly	
545 550 555 560	
tac gaa ttg ggc gcg tcc tac cgc acc ggc ggc ctg act gcc aaa gtc	1728
Tyr Glu Leu Gly Ala Ser Tyr Arg Thr Gly Gly Leu Thr Ala Lys Val	
565 570 575	
ggc gta agc cac agc aaa ccg cgc ttt tac gat acg cac aaa gac aag	1776
Gly Val Ser His Ser Lys Pro Arg Phe Tyr Asp Thr His Lys Asp Lys	
580 585 590	

ctg ttg agc gcg aat cct gaa ttt ggc gca caa gtc ggc cgc act tgg 1824
 Leu Leu Ser Ala Asn Pro Glu Phe Gly Ala Gln Val Gly Arg Thr Trp
 595 600 605

acg gct tcc ctt gcc tac cgc ttc caa aac ccg aat ctg gaa atc ggc 1872
 Thr Ala Ser Leu Ala Tyr Arg Phe Gln Asn Pro Asn Leu Glu Ile Gly
 610 615 620

tgg cgc ggc cgt tat gtt caa aaa gcc gtg ggt tcg ata ttg gtg gca 1920
 Trp Arg Gly Arg Tyr Val Gln Lys Ala Val Gly Ser Ile Leu Val Ala
 625 630 635 640

ggt caa aaa gac cgc aac ggc aaa ttg gaa aac gtt gta cgc aaa ggt 1968
 Gly Gln Lys Asp Arg Asn Gly Lys Leu Glu Asn Val Val Arg Lys Gly
 645 650 655

ttc ggt gtg aac gat gtc ttc gcc aac tgg aaa ccg ctg ggc aaa gac 2016
 Phe Gly Val Asn Asp Val Phe Ala Asn Trp Lys Pro Leu Gly Lys Asp
 660 665 670

acg ctc aat gtt aat ctt tcg gtt aac aac gtg ttc aac acg ttc tac 2064
 Thr Leu Asn Val Asn Leu Ser Val Asn Asn Val Phe Asn Thr Phe Tyr
 675 680 685

tat ccg cac agc caa cga tgg acc aat acc ctg ccg ggc gtg gga cgt 2112
 Tyr Pro His Ser Gln Arg Trp Thr Asn Thr Leu Pro Gly Val Gly Arg
 690 695 700

gat gta cgc ttg ggc gtg aac tac aag ttc taa 2145
 Asp Val Arg Leu Gly Val Asn Tyr Lys Phe
 705 710 715

<210> 196

<211> 714

<212> PRT

<213> Neisseria meningitidis

<400> 196

Met Asn Thr Pro Leu Phe Arg Leu Ser Leu Leu Ser Leu Thr Leu Ala
 1 5 10 15

Ala Gly Phe Ala His Ala Ala Glu Asn Asn Ala Lys Val Val Leu Asp
 20 25 30

Thr Val Thr Val Lys Gly Asp Arg Gln Gly Ser Lys Ile Arg Thr Asn
 35 40 45

Ile Val Thr Leu Gln Gln Lys Asp Glu Ser Thr Ala Thr Asp Met Arg
 50 55 60

Glu Leu Leu Lys Glu Glu Pro Ser Ile Asp Phe Gly Gly Gly Asn Gly
 65 70 75 80

Thr Ser Gln Phe Leu Thr Leu Arg Gly Met Gly Gln Asn Ser Val Asp
 85 90 95

Ile Lys Val Asp Asn Ala Tyr Ser Asp Ser Gln Ile Leu Tyr His Gln
 100 105 110

Gly Arg Phe Ile Val Asp Pro Ala Leu Val Lys Val Val Ser Val Gln
 115 120 125

Lys Gly Ala Gly Ser Ala Ser Ala Gly Ile Gly Ala Thr Asn Gly Ala
 130 135 140

Ile Ile Thr Lys Thr Val Asp Ala Gln Asp Leu Leu Lys Gly Leu Asp
 145 150 155 160

Lys Asn Trp Gly Val Arg Leu Asn Ser Gly Phe Ala Ser Asn Glu Gly
 165 170 175

Val Ser Tyr Gly Ala Ser Val Phe Gly Lys Glu Gly Asn Phe Asp Gly
 180 185 190

Leu Phe Ser Tyr Asn Arg Asn Asn Glu Lys Asp Tyr Glu Ala Gly Lys
 195 200 205

Gly Phe Arg Asn Asn Phe Asn Gly Gly Lys Thr Val Pro Tyr Ser Ala
 210 215 220

Leu Asp Lys Arg Ser Tyr Leu Ala Lys Ile Gly Thr Ser Phe Gly Asp
 225 230 235 240

Gly Asp His Arg Ile Val Leu Ser His Met Lys Asp Gln His Arg Gly
 245 250 255

Ile Arg Thr Val Arg Glu Glu Phe Thr Val Gly Gly Asp Lys Glu Arg
 260 265 270

Ile Ser Met Glu Arg Gln Ala Pro Ala Tyr Arg Glu Thr Thr Gln Ser
 275 280 285

Asn Thr Asn Leu Ala Tyr Thr Gly Lys Asn Leu Gly Phe Val Glu Lys
 290 295 300

Leu Asp Ala Asn Ala Tyr Val Leu Glu Lys Glu Arg Tyr Ser Ala Asp
 305 310 315 320
 Asp Ser Gly Thr Gly Tyr Ala Gly Asn Val Lys Gly Pro Asn His Thr
 325 330 335
 Gln Ile Thr Thr Arg Gly Met Asn Phe Asn Phe Asp Ser Arg Leu Ala
 340 345 350
 Glu Gln Thr Leu Leu Lys Tyr Gly Ile Asn Tyr Arg His Gln Glu Ile
 355 360 365
 Lys Pro Gln Ala Phe Leu Asn Ser Gln Phe Lys Ile Glu Asp Lys Glu
 370 375 380
 Lys Ala Thr Asp Glu Glu Lys Asn Lys Asn Arg Glu Asn Glu Lys Ile
 385 390 395 400
 Ala Lys Ala Tyr Arg Leu Thr Asn Pro Thr Lys Thr Asp Thr Gly Ala
 405 410 415
 Tyr Ile Glu Ala Ile His Glu Ile Asp Gly Phe Thr Leu Thr Gly Gly
 420 425 430
 Leu Arg Tyr Asp Arg Phe Lys Val Lys Thr His Asp Gly Lys Thr Val
 435 440 445
 Ser Ser Asn Asn Leu Asn Pro Ser Phe Gly Val Ile Trp Gln Pro His
 450 455 460
 Glu His Trp Ser Phe Ser Ala Ser His Asn Tyr Ala Ser Arg Ser Pro
 465 470 475 480
 Arg Leu Tyr Asp Ala Leu Gln Thr His Gly Lys Arg Gly Ile Ile Ser
 485 490 495
 Ile Ala Asp Gly Thr Lys Ala Glu Arg Ala Arg Asn Thr Glu Ile Gly
 500 505 510
 Phe Asn Tyr Asn Asp Gly Thr Phe Ala Ala Asn Gly Ser Tyr Phe Trp
 515 520 525
 Gln Thr Ile Lys Asp Ala Leu Ala Asn Pro Gln Asn Arg His Asp Ser
 530 535 540
 Val Ala Val Arg Glu Ala Val Asn Ala Gly Tyr Ile Lys Asn His Gly
 545 550 555 560

Tyr Glu Leu Gly Ala Ser Tyr Arg Thr Gly Gly Leu Thr Ala Lys Val
 565 570 575

Gly Val Ser His Ser Lys Pro Arg Phe Tyr Asp Thr His Lys Asp Lys
 580 585 590

Leu Leu Ser Ala Asn Pro Glu Phe Gly Ala Gln Val Gly Arg Thr Trp
 595 600 605

Thr Ala Ser Leu Ala Tyr Arg Phe Gln Asn Pro Asn Leu Glu Ile Gly
 610 615 620

Trp Arg Gly Arg Tyr Val Gln Lys Ala Val Gly Ser Ile Leu Val Ala
 625 630 635 640

Gly Gln Lys Asp Arg Asn Gly Lys Leu Glu Asn Val Val Arg Lys Gly
 645 650 655

Phe Gly Val Asn Asp Val Phe Ala Asn Trp Lys Pro Leu Gly Lys Asp
 660 665 670

Thr Leu Asn Val Asn Leu Ser Val Asn Asn Val Phe Asn Thr Phe Tyr
 675 680 685

Tyr Pro His Ser Gln Arg Trp Thr Asn Thr Leu Pro Gly Val Gly Arg
 690 695 700

Asp Val Arg Leu Gly Val Asn Tyr Lys Phe
 705 710

<210> 197

<211> 2625

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(2625)

<400> 197

atg aaa acc tcc gaa ctg cgc caa aaa ttc cta aaa ttt ttt gaa acc 48
 Met Lys Thr Ser Glu Leu Arg Gln Lys Phe Leu Lys Phe Phe Glu Thr
 1 5 10 15

aaa ggc cac acc gtc gtc cgc tct tcc agc ctc gtg ccg cac gac gac 96

Lys Gly His Thr Val Val Arg Ser Ser Ser Leu Val Pro His Asp Asp	
20 25 30	
ccg acc ctg ctg ttt acc aac gcg ggc atg aac cag ttt aaa gac gta	144
Pro Thr Leu Leu Phe Thr Asn Ala Gly Met Asn Gln Phe Lys Asp Val	
35 40 45	
ttc tta ggt ttc gac aaa cgc ccg tac agc cgc gcc acc acc gcg caa	192
Phe Leu Gly Phe Asp Lys Arg Pro Tyr Ser Arg Ala Thr Thr Ala Gln	
50 55 60	
aaa tgc gta cgc gca ggc ggc aaa cac aac gac ttg gaa aac gtc ggc	240
Lys Cys Val Arg Ala Gly Gly Lys His Asn Asp Leu Glu Asn Val Gly	
65 70 75 80	
tac acc gcc cgc cac cac acc ttc ttt gaa atg atg ggc aac ttc tcc	288
Tyr Thr Ala Arg His His Thr Phe Phe Glu Met Met Gly Asn Phe Ser	
85 90 95	
ttc ggc gac tac ttc aaa cgc gac gcc atc cac ttc gct tgg gaa ttt	336
Phe Gly Asp Tyr Phe Lys Arg Asp Ala Ile His Phe Ala Trp Glu Phe	
100 105 110	
ctg act tcc ccc gaa tgg ctc aac atc cct aaa gac aaa ctg ttg gcg	384
Leu Thr Ser Pro Glu Trp Leu Asn Ile Pro Lys Asp Lys Leu Leu Ala	
115 120 125	
acc gtt tac gcg gaa gac gac gaa gcc tac aac atc tgg ttg aac gaa	432
Thr Val Tyr Ala Glu Asp Asp Glu Ala Tyr Asn Ile Trp Leu Asn Glu	
130 135 140	
atc ggt atg ccg tcc gag cgc atc gtc cgc atc ggc gac aac aaa ggc	480
Ile Gly Met Pro Ser Glu Arg Ile Val Arg Ile Gly Asp Asn Lys Gly	
145 150 155 160	
gcg aaa tac gca tcc gac aac ttc tgg caa atg ggc gac acc ggc cct	528
Ala Lys Tyr Ala Ser Asp Asn Phe Trp Gln Met Gly Asp Thr Gly Pro	
165 170 175	
tgc ggc ccc tgc tcc gaa att ttc tac gac cac ggc gaa gaa atc tgg	576
Cys Gly Pro Cys Ser Glu Ile Phe Tyr Asp His Gly Glu Glu Ile Trp	
180 185 190	
ggc ggc att ccc ggc agt ccc gaa gaa gac ggc gac cgc tgg atc gaa	624
Gly Gly Ile Pro Gly Ser Pro Glu Glu Asp Gly Asp Arg Trp Ile Glu	
195 200 205	
att tgg aac tgc gta ttt atg cag ttc aac cgc gac gaa caa ggc aat	672

Ile Trp Asn Cys Val Phe Met Gln Phe Asn Arg Asp Glu Gln Gly Asn	
210	220
atg aac ccg ctt ccc aaa cct tcc gtc gat acc ggt atg ggc ttg gaa	720
Met Asn Pro Leu Pro Lys Pro Ser Val Asp Thr Gly Met Gly Leu Glu	
225	240
cgc ata gcc gcc gtc atg cag cat gtt cac agc aac tac gaa atc gac	768
Arg Ile Ala Ala Val Met Gln His Val His Ser Asn Tyr Glu Ile Asp	
245	255
ttg ttc caa gac ctg ctc aaa gcc gtt gcc cgc gaa acc ggc gcg ccg	816
Leu Phe Gln Asp Leu Leu Lys Ala Val Ala Arg Glu Thr Gly Ala Pro	
260	270
ttc aga atg gaa gaa ccc agc ctg aaa gtc atc gcc gac cac atc cgc	864
Phe Arg Met Glu Glu Pro Ser Leu Lys Val Ile Ala Asp His Ile Arg	
275	285
tcc tgc tcg ttc ctg att gca gac ggc gtc ttg cct tcc aac gaa ggc	912
Ser Cys Ser Phe Leu Ile Ala Asp Gly Val Leu Pro Ser Asn Glu Gly	
290	300
cgc ggc tac gta ttg cgc cgc att atc cgc cgc gcc gtg cgc cac ggt	960
Arg Gly Tyr Val Leu Arg Arg Ile Ile Arg Arg Ala Val Arg His Gly	
305	320
tac aaa ctg ggt caa agc aaa ccg ttc ttc cac aaa ctc gtt gcc gat	1008
Tyr Lys Leu Gly Gln Ser Lys Pro Phe Phe His Lys Leu Val Ala Asp	
325	335
ttg gtc aaa gag atg ggc ggt gcc tac cct gaa ttg aaa gaa aaa caa	1056
Leu Val Lys Glu Met Gly Gly Ala Tyr Pro Glu Leu Lys Glu Lys Gln	
340	350
gcc caa atc gaa gaa gca ttg aaa aac gaa gaa agc cgt ttt gcc caa	1104
Ala Gln Ile Glu Glu Ala Leu Lys Asn Glu Glu Ser Arg Phe Ala Gln	
355	365
acg ctg gaa acc ggt atg gct ttg ttg gaa aac gcg ctg gtc aaa ggc	1152
Thr Leu Glu Thr Gly Met Ala Leu Leu Glu Asn Ala Leu Val Lys Gly	
370	380
ggc aaa aca ctc ggc ggc gaa atc atc ttc aaa ctc tac gat acc tac	1200
Gly Lys Thr Leu Gly Gly Glu Ile Ile Phe Lys Leu Tyr Asp Thr Tyr	
385	400
ggg ttc cca tac gac ttg act gcc gac atc tgc cgc gaa cgc aat atc	1248

Gly Phe Pro Tyr Asp Leu Thr Ala Asp Ile Cys Arg Glu Arg Asn Ile	
405 410 415	
gaa ccg gac gaa gca ggc ttc gag cgc gaa atg gaa gcc caa cgc gca	1296
Glu Pro Asp Glu Ala Gly Phe Glu Arg Glu Met Glu Ala Gln Arg Ala	
420 425 430	
cgc gca cgc gcc gcc caa agc ttc aaa gcc aac gcc caa ctg cct tat	1344
Arg Ala Arg Ala Ala Gln Ser Phe Lys Ala Asn Ala Gln Leu Pro Tyr	
435 440 445	
gac ggt caa gac acc gag ttt aaa ggt tat agc gaa cgc caa acc gaa	1392
Asp Gly Gln Asp Thr Glu Phe Lys Gly Tyr Ser Glu Arg Gln Thr Glu	
450 455 460	
tcc aaa gtc ctc gcc ctc tac aaa gac ggc gag caa gtc aac gaa ttg	1440
Ser Lys Val Leu Ala Leu Tyr Lys Asp Gly Glu Gln Val Asn Glu Leu	
465 470 475 480	
aac gaa ggc gac agc ggc gca gtc gtc atc gac ttt acc ccg ttc tat	1488
Asn Glu Gly Asp Ser Gly Ala Val Val Ile Asp Phe Thr Pro Phe Tyr	
485 490 495	
gca gaa tcc ggc ggc caa gtc ggc gat gtc ggc tat atc ttc tca ggc	1536
Ala Glu Ser Gly Gly Gln Val Gly Asp Val Gly Tyr Ile Phe Ser Gly	
500 505 510	
gaa aac cgc ttt gaa gta cgc gat acc caa aaa atc aaa gcg gcc gta	1584
Glu Asn Arg Phe Glu Val Arg Asp Thr Gln Lys Ile Lys Ala Ala Val	
515 520 525	
ttc ggt caa ttc ggc gta caa act tca ggc cgt ctg aaa gtc ggc gac	1632
Phe Gly Gln Phe Gly Val Gln Thr Ser Gly Arg Leu Lys Val Gly Asp	
530 535 540	
agc gtt acc gcc aaa gtg gac gac gaa atc cgc aat gcc aat atg cgc	1680
Ser Val Thr Ala Lys Val Asp Asp Glu Ile Arg Asn Ala Asn Met Arg	
545 550 555 560	
aac cac agc gca acc cac ttg atg cac aaa gcc ctg cgc gat gta ttg	1728
Asn His Ser Ala Thr His Leu Met His Lys Ala Leu Arg Asp Val Leu	
565 570 575	
ggc aga cac gtc gaa caa aaa ggc tct ttg gtt acc gcc gaa tcc acc	1776
Gly Arg His Val Glu Gln Lys Gly Ser Leu Val Thr Ala Glu Ser Thr	
580 585 590	
cgt ttc gac att tcc cat ccc caa gcg gta act gcc gaa gaa att gcc	1824

Arg Phe Asp Ile Ser His Pro Gln Ala Val Thr Ala Glu Glu Ile-Ala	
595 600 605	
gaa gta gaa cgc cgc gtc aac gaa gcc att ttg gcg aac gtt gcc gtc	1872
Glu Val Glu Arg Arg Val Asn Glu Ala Ile Leu Ala Asn Val Ala Val	
610 615 620	
aat gca gcc att atg agc atg gaa gac gcg caa aaa acc ggc gcg atg	1920
Asn Ala Ala Ile Met Ser Met Glu Asp Ala Gln Lys Thr Gly Ala Met	
625 630 635 640	
atg ctc ttc ggc gaa aaa tac ggc gaa gaa gtg cgc gta ctg caa atg	1968
Met Leu Phe Gly Glu Lys Tyr Gly Glu Glu Val Arg Val Leu Gln Met	
645 650 655	
ggc ggt ttc tct acc gaa ttg tgc ggc ggc aca cac gtt tca cgc acc	2016
Gly Gly Phe Ser Thr Glu Leu Cys Gly Gly Thr His Val Ser Arg Thr	
660 665 670	
ggc gac atc ggc ctc ttc aaa atc atc agc gaa ggc ggt att gcc gca	2064
Gly Asp Ile Gly Leu Phe Lys Ile Ile Ser Glu Gly Gly Ile Ala Ala	
675 680 685	
ggc gtg cgc cgt atc gaa gcc atc acc ggc ctg aac gca ctc aaa tgg	2112
Gly Val Arg Arg Ile Glu Ala Ile Thr Gly Leu Asn Ala Leu Lys Trp	
690 695 700	
gcg caa gag caa gag cgt ttg gtg aaa gac att att gcc gaa acc aaa	2160
Ala Gln Glu Gln Glu Arg Leu Val Lys Asp Ile Ile Ala Glu Thr Lys	
705 710 715 720	
gcc caa acc gaa aaa gac gta ctg gca aaa atc caa gca ggc gcg gca	2208
Ala Gln Thr Glu Lys Asp Val Leu Ala Lys Ile Gln Ala Gly Ala Ala	
725 730 735	
cac gcc aaa gca ttg gaa aaa gaa ttg gca cgc gcc aaa gcc gaa ctc	2256
His Ala Lys Ala Leu Glu Lys Glu Leu Ala Arg Ala Lys Ala Glu Leu	
740 745 750	
gcc gtc cac gca ggc gcc aaa ctc ttg gac gat gca aaa gac ttg ggc	2304
Ala Val His Ala Gly Ala Lys Leu Leu Asp Asp Ala Lys Asp Leu Gly	
755 760 765	
gca gcc aaa ctc gtt gcc gcc caa atc gaa gcc gac gca gcc gcc ctg	2352
Ala Ala Lys Leu Val Ala Ala Gln Ile Glu Ala Asp Ala Ala Ala Leu	
770 775 780	
cgc gaa atc gtt acc gat tta acc ggt aaa tcc gac aac gcc gtg att	2400

Arg Glu Ile Val Thr Asp Leu Thr Gly Lys Ser Asp Asn Ala Val Ile
 785 790 795 800

ctt tta gcg gca gta aac gac ggc aaa gtc tcc ctg tgc gcc ggc gta 2448
 Leu Leu Ala Ala Val Asn Asp Gly Lys Val Ser Leu Cys Ala Gly Val
 805 810 815

tcc aaa ccg ttg acc ggc aaa gtg aaa gca ggc gat ctg gtt aaa ttt 2496
 Ser Lys Pro Leu Thr Gly Lys Val Lys Ala Gly Asp Leu Val Lys Phe
 820 825 830

gca gcc gaa caa gtc ggc ggc aaa ggc ggc ggc aga cca gat ttg gcg 2544
 Ala Ala Glu Gln Val Gly Gly Lys Gly Gly Gly Arg Pro Asp Leu Ala
 835 840 845

caa gcc ggc ggc acg gat gcc gac aaa ttg ccc gcc gtg ttg gat agc 2592
 Gln Ala Gly Gly Thr Asp Ala Asp Lys Leu Pro Ala Val Leu Asp Ser
 850 855 860

gtg aaa gac tgg gtc ggc gcg aag ctg gtt tga 2625
 Val Lys Asp Trp Val Gly Ala Lys Leu Val
 865 870 875

<210> 198

<211> 874

<212> PRT

<213> Neisseria meningitidis

<400> 198

Met Lys Thr Ser Glu Leu Arg Gln Lys Phe Leu Lys Phe Phe Glu Thr
 1 5 10 15

Lys Gly His Thr Val Val Arg Ser Ser Ser Leu Val Pro His Asp Asp
 20 25 30

Pro Thr Leu Leu Phe Thr Asn Ala Gly Met Asn Gln Phe Lys Asp Val
 35 40 45

Phe Leu Gly Phe Asp Lys Arg Pro Tyr Ser Arg Ala Thr Thr Ala Gln
 50 55 60

Lys Cys Val Arg Ala Gly Gly Lys His Asn Asp Leu Glu Asn Val Gly
 65 70 75 80

Tyr Thr Ala Arg His His Thr Phe Phe Glu Met Met Gly Asn Phe Ser
 85 90 95

Phe Gly Asp Tyr Phe Lys Arg Asp Ala Ile His Phe Ala Trp Glu Phe
 100 105 110

Leu Thr Ser Pro Glu Trp Leu Asn Ile Pro Lys Asp Lys Leu Leu Ala
 115 120 125

Thr Val Tyr Ala Glu Asp Asp Glu Ala Tyr Asn Ile Trp Leu Asn Glu
 130 135 140

Ile Gly Met Pro Ser Glu Arg Ile Val Arg Ile Gly Asp Asn Lys Gly
 145 150 155 160

Ala Lys Tyr Ala Ser Asp Asn Phe Trp Gln Met Gly Asp Thr Gly Pro
 165 170 175

Cys Gly Pro Cys Ser Glu Ile Phe Tyr Asp His Gly Glu Glu Ile Trp
 180 185 190

Gly Gly Ile Pro Gly Ser Pro Glu Glu Asp Gly Asp Arg Trp Ile Glu
 195 200 205

Ile Trp Asn Cys Val Phe Met Gln Phe Asn Arg Asp Glu Gln Gly Asn
 210 215 220

Met Asn Pro Leu Pro Lys Pro Ser Val Asp Thr Gly Met Gly Leu Glu
 225 230 235 240

Arg Ile Ala Ala Val Met Gln His Val His Ser Asn Tyr Glu Ile Asp
 245 250 255

Leu Phe Gln Asp Leu Leu Lys Ala Val Ala Arg Glu Thr Gly Ala Pro
 260 265 270

Phe Arg Met Glu Glu Pro Ser Leu Lys Val Ile Ala Asp His Ile Arg
 275 280 285

Ser Cys Ser Phe Leu Ile Ala Asp Gly Val Leu Pro Ser Asn Glu Gly
 290 295 300

Arg Gly Tyr Val Leu Arg Arg Ile Ile Arg Arg Ala Val Arg His Gly
 305 310 315 320

Tyr Lys Leu Gly Gln Ser Lys Pro Phe Phe His Lys Leu Val Ala Asp
 325 330 335

Leu Val Lys Glu Met Gly Gly Ala Tyr Pro Glu Leu Lys Glu Lys Gln
 340 345 350

Ala Gln Ile Glu Glu Ala Leu Lys Asn Glu Glu Ser Arg Phe Ala Gln
 355 360 365
 Thr Leu Glu Thr Gly Met Ala Leu Leu Glu Asn Ala Leu Val Lys Gly
 370 375 380
 Gly Lys Thr Leu Gly Gly Glu Ile Ile Phe Lys Leu Tyr Asp Thr Tyr
 385 390 395 400
 Gly Phe Pro Tyr Asp Leu Thr Ala Asp Ile Cys Arg Glu Arg Asn Ile
 405 410 415
 Glu Pro Asp Glu Ala Gly Phe Glu Arg Glu Met Glu Ala Gln Arg Ala
 420 425 430
 Arg Ala Arg Ala Ala Gln Ser Phe Lys Ala Asn Ala Gln Leu Pro Tyr
 435 440 445
 Asp Gly Gln Asp Thr Glu Phe Lys Gly Tyr Ser Glu Arg Gln Thr Glu
 450 455 460
 Ser Lys Val Leu Ala Leu Tyr Lys Asp Gly Glu Gln Val Asn Glu Leu
 465 470 475 480
 Asn Glu Gly Asp Ser Gly Ala Val Val Ile Asp Phe Thr Pro Phe Tyr
 485 490 495
 Ala Glu Ser Gly Gly Gln Val Gly Asp Val Gly Tyr Ile Phe Ser Gly
 500 505 510
 Glu Asn Arg Phe Glu Val Arg Asp Thr Gln Lys Ile Lys Ala Ala Val
 515 520 525
 Phe Gly Gln Phe Gly Val Gln Thr Ser Gly Arg Leu Lys Val Gly Asp
 530 535 540
 Ser Val Thr Ala Lys Val Asp Asp Glu Ile Arg Asn Ala Asn Met Arg
 545 550 555 560
 Asn His Ser Ala Thr His Leu Met His Lys Ala Leu Arg Asp Val Leu
 565 570 575
 Gly Arg His Val Glu Gln Lys Gly Ser Leu Val Thr Ala Glu Ser Thr
 580 585 590
 Arg Phe Asp Ile Ser His Pro Gln Ala Val Thr Ala Glu Glu Ile Ala
 595 600 605

Glu Val Glu Arg Arg Val Asn Glu Ala Ile Leu Ala Asn Val Ala Val
 610 615 620

Asn Ala Ala Ile Met Ser Met Glu Asp Ala Gln Lys Thr Gly Ala Met
 625 630 635 640

Met Leu Phe Gly Glu Lys Tyr Gly Glu Glu Val Arg Val Leu Gln Met
 645 650 655

Gly Gly Phe Ser Thr Glu Leu Cys Gly Gly Thr His Val Ser Arg Thr
 660 665 670

Gly Asp Ile Gly Leu Phe Lys Ile Ile Ser Glu Gly Gly Ile Ala Ala
 675 680 685

Gly Val Arg Arg Ile Glu Ala Ile Thr Gly Leu Asn Ala Leu Lys Trp
 690 695 700

Ala Gln Glu Gln Glu Arg Leu Val Lys Asp Ile Ile Ala Glu Thr Lys
 705 710 715 720

Ala Gln Thr Glu Lys Asp Val Leu Ala Lys Ile Gln Ala Gly Ala Ala
 725 730 735

His Ala Lys Ala Leu Glu Lys Glu Leu Ala Arg Ala Lys Ala Glu Leu
 740 745 750

Ala Val His Ala Gly Ala Lys Leu Leu Asp Asp Ala Lys Asp Leu Gly
 755 760 765

Ala Ala Lys Leu Val Ala Ala Gln Ile Glu Ala Asp Ala Ala Ala Leu
 770 775 780

Arg Glu Ile Val Thr Asp Leu Thr Gly Lys Ser Asp Asn Ala Val Ile
 785 790 795 800

Leu Leu Ala Ala Val Asn Asp Gly Lys Val Ser Leu Cys Ala Gly Val
 805 810 815

Ser Lys Pro Leu Thr Gly Lys Val Lys Ala Gly Asp Leu Val Lys Phe
 820 825 830

Ala Ala Glu Gln Val Gly Gly Lys Gly Gly Gly Arg Pro Asp Leu Ala
 835 840 845

Gln Ala Gly Gly Thr Asp Ala Asp Lys Leu Pro Ala Val Leu Asp Ser
 850 855 860

Val Lys Asp Trp Val Gly Ala Lys Leu Val
865 870

<210> 199
<211> 207
<212> DNA
<213> Neisseria meningitidis

<220>
<221> CDS
<222> (1)..(207)

<400> 199
gtg caa aga ggc cgt ctg aaa ggt ctc gtt tgc cgt agg ttg ggt cgc 48
Val Gln Arg Gly Arg Leu Lys Gly Leu Val Cys Arg Arg Leu Gly Arg
1 5 10 15
gac cca aca aat ttt gtg aag tat aaa aat gtt ggt cat gac cca acc 96
Asp Pro Thr Asn Phe Val Lys Tyr Lys Asn Val Gly His Asp Pro Thr
20 25 30
tac ctg cct ttt tgt aca aag agg cta tct gaa agg cct tgt ttg ccg 144
Tyr Leu Pro Phe Cys Thr Lys Arg Leu Ser Glu Arg Pro Cys Leu Pro
35 40 45
tat ggt ggg tcg cga ccc agc aga ttt tta tta ggg tat gac cca agc 192
Tyr Gly Gly Ser Arg Pro Ser Arg Phe Leu Leu Gly Tyr Asp Pro Ser
50 55 60
tac ttg cta cga taa 207
Tyr Leu Leu Arg
65

<210> 200
<211> 68
<212> PRT
<213> Neisseria meningitidis

<400> 200
Val Gln Arg Gly Arg Leu Lys Gly Leu Val Cys Arg Arg Leu Gly Arg
1 5 10 15
Asp Pro Thr Asn Phe Val Lys Tyr Lys Asn Val Gly His Asp Pro Thr
20 25 30

Tyr Leu Pro Phe Cys Thr Lys Arg Leu Ser Glu Arg Pro Cys Leu Pro
 35 40 45

Tyr Gly Gly Ser Arg Pro Ser Arg Phe Leu Leu Gly Tyr Asp Pro Ser
 50 55 60

Tyr Leu Leu Arg
 65

<210> 201

<211> 2277

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(2277)

<400> 201

atg gca caa act aca ctc aaa ccc att gtt tta tca att ctt tta atc	48
Met Ala Gln Thr Thr Leu Lys Pro Ile Val Leu Ser Ile Leu Leu Ile	
1 5 10 15	
aac aca ccc ctc ctc gcc caa gcg cat gaa act gag caa tcg gtg gat	96
Asn Thr Pro Leu Leu Ala Gln Ala His Glu Thr Glu Gln Ser Val Asp	
20 25 30	
ttg gaa acg gtc agc gtc gtc ggc aaa agc cgt ccg cgc gcc acg tcg	144
Leu Glu Thr Val Ser Val Val Gly Lys Ser Arg Pro Arg Ala Thr Ser	
35 40 45	
ggg ctg ttg cac act tcg acc gcc tcc gac aaa atc atc tcc ggc gat	192
Gly Leu Leu His Thr Ser Thr Ala Ser Asp Lys Ile Ile Ser Gly Asp	
50 55 60	
acc ttg cgc caa aaa gcc gtc aac ttg ggc gac gct tta gac ggc gta	240
Thr Leu Arg Gln Lys Ala Val Asn Leu Gly Asp Ala Leu Asp Gly Val	
65 70 75 80	
ccg ggc atc cac gct tcg caa tac ggc ggc ggc gcg tct gct ccc gtc	288
Pro Gly Ile His Ala Ser Gln Tyr Gly Gly Gly Ala Ser Ala Pro Val	
85 90 95	
att cgc ggt caa aca ggc agg cgg att aaa gtg ttg aac cat cac ggc	336
Ile Arg Gly Gln Thr Gly Arg Arg Ile Lys Val Leu Asn His His Gly	
100 105 110	

gaa aca ggc gat atg gcg gat ttt tcg ccc gat cac gcc att atg gta 384
 Glu Thr Gly Asp Met Ala Asp Phe Ser Pro Asp His Ala Ile Met Val
 115 120 125

gat acc gcc ttg tcg caa cag gtc gaa atc ctg cgc ggg ccg gtt acg 432
 Asp Thr Ala Leu Ser Gln Gln Val Glu Ile Leu Arg Gly Pro Val Thr
 130 135 140

ctc ttg tac agc tcg ggc aat gtg gcg ggg ctg gtc gat gtt gcc gat 480
 Leu Leu Tyr Ser Ser Gly Asn Val Ala Gly Leu Val Asp Val Ala Asp
 145 150 155 160

ggc aaa atc ccc gaa aaa atg cct gaa aac ggc gta tcg ggc gaa ctc 528
 Gly Lys Ile Pro Glu Lys Met Pro Glu Asn Gly Val Ser Gly Glu Leu
 165 170 175

gga ttg cgt ttg agc agc ggc aat ctg gaa aaa ctc acg tcc ggc ggc 576
 Gly Leu Arg Leu Ser Ser Gly Asn Leu Glu Lys Leu Thr Ser Gly Gly
 180 185 190

atc aat atc ggt ttg ggc aaa aac ttt gta ttg cac acg gaa ggg ctg 624
 Ile Asn Ile Gly Leu Gly Lys Asn Phe Val Leu His Thr Glu Gly Leu
 195 200 205

tac cgc aaa tcg ggg gat tac gcc gta ccg cgt tac cgc aat ctg aaa 672
 Tyr Arg Lys Ser Gly Asp Tyr Ala Val Pro Arg Tyr Arg Asn Leu Lys
 210 215 220

cgc ctg ccc gac agc cac gcc gat tcg caa acg ggc agc atc ggg ctg 720
 Arg Leu Pro Asp Ser His Ala Asp Ser Gln Thr Gly Ser Ile Gly Leu
 225 230 235 240

tct tgg gtt ggc gaa aaa ggt ttt atc ggc gta gcg tac agc gac cgt 768
 Ser Trp Val Gly Glu Lys Gly Phe Ile Gly Val Ala Tyr Ser Asp Arg
 245 250 255

cgc gac caa tat ggt ctg cct gcc cac agc cac gaa tac gat gat tgc 816
 Arg Asp Gln Tyr Gly Leu Pro Ala His Ser His Glu Tyr Asp Asp Cys
 260 265 270

cac gcc gac atc atc tgg caa aag agc ttg att aac aaa cgc tat tta 864
 His Ala Asp Ile Ile Trp Gln Lys Ser Leu Ile Asn Lys Arg Tyr Leu
 275 280 285

cag ctt tat ccg cac ctg ttg acc gaa gaa gac atc gat tac gac aat 912
 Gln Leu Tyr Pro His Leu Leu Thr Glu Glu Asp Ile Asp Tyr Asp Asn
 290 295 300

396

cca caa cac aaa ctc agc ctg acc gcc tcc cat cag gaa cgc ctg ccg	1536
Pro Gln His Lys Leu Ser Leu Thr Ala Ser His Gln Glu Arg Leu Pro	
500 505 510	
tca acg caa gag ctg tac gca cac ggc aaa cac gtc gcc acc aac acc	1584
Ser Thr Gln Glu Leu Tyr Ala His Gly Lys His Val Ala Thr Asn Thr	
515 520 525	
ttt gaa gtc ggc aac aaa cac ctc aac aaa gag cgt tcc aac aat atc	1632
Phe Glu Val Gly Asn Lys His Leu Asn Lys Glu Arg Ser Asn Asn Ile	
530 535 540	
gaa ctc gcg ctg ggc tac gaa ggc gac cgc tgg caa tac aat ctg gca	1680
Glu Leu Ala Leu Gly Tyr Glu Gly Asp Arg Trp Gln Tyr Asn Leu Ala	
545 550 555 560	
ctc tac cgc aac cgc ttc ggt aac tac att tac gcc caa acc tta aac	1728
Leu Tyr Arg Asn Arg Phe Gly Asn Tyr Ile Tyr Ala Gln Thr Leu Asn	
565 570 575	
gac gga cgc ggc ccc aaa tcc atc gaa gac gac agc gaa atg aag ctc	1776
Asp Gly Arg Gly Pro Lys Ser Ile Glu Asp Asp Ser Glu Met Lys Leu	
580 585 590	
gtg cgc tac aac caa tcc ggc gcc gac ttc tac ggc gcg gaa ggc gaa	1824
Val Arg Tyr Asn Gln Ser Gly Ala Asp Phe Tyr Gly Ala Glu Gly Glu	
595 600 605	
atc tac ttc aaa ccg aca ccg cgc tac cgc atc ggc gtt tcc ggc gac	1872
Ile Tyr Phe Lys Pro Thr Pro Arg Tyr Arg Ile Gly Val Ser Gly Asp	
610 615 620	
tat gta cga ggc cgt ctg aaa aac ctg cct tcc cta ccc ggc aga gaa	1920
Tyr Val Arg Gly Arg Leu Lys Asn Leu Pro Ser Leu Pro Gly Arg Glu	
625 630 635 640	
gat gcc tac ggc aac cgt cct ttc atc gca cag gac gac caa aat gcc	1968
Asp Ala Tyr Gly Asn Arg Pro Phe Ile Ala Gln Asp Asp Gln Asn Ala	
645 650 655	
ccc cgt gtt ccg gct gcg cgc ctc ggc ttc cac ctg aaa gcc tcg ctg	2016
Pro Arg Val Pro Ala Ala Arg Leu Gly Phe His Leu Lys Ala Ser Leu	
660 665 670	
acc gac cgt atc gat gcc aat ttg gac tac tac cgc gtg ttc gcc caa	2064
Thr Asp Arg Ile Asp Ala Asn Leu Asp Tyr Tyr Arg Val Phe Ala Gln	
675 680 685	

aac aaa ctc gcc cgc tac gaa acg cgc acg ccc gga cac cat atg ctc 2112
 Asn Lys Leu Ala Arg Tyr Glu Thr Arg Thr Pro Gly His His Met Leu
 690 695 700

aac ctc ggc gca aac tac cgc cgc aat acg cgc tat ggc gag tgg aat 2160
 Asn Leu Gly Ala Asn Tyr Arg Arg Asn Thr Arg Tyr Gly Glu Trp Asn
 705 710 715 720

tgg tac gtc aaa gcc gac aac ctg ctc aac caa tcc gtt tac gcc cac 2208
 Trp Tyr Val Lys Ala Asp Asn Leu Leu Asn Gln Ser Val Tyr Ala His
 725 730 735

agc agc ttt ctc tct gat acg ccg caa atg ggc cgc agc ttt acc ggc 2256
 Ser Ser Phe Leu Ser Asp Thr Pro Gln Met Gly Arg Ser Phe Thr Gly
 740 745 750

ggc gtg aac gtg aag ttt taa 2277
 Gly Val Asn Val Lys Phe
 755

<210> 202

<211> 758

<212> PRT

<213> Neisseria meningitidis

<400> 202

Met Ala Gln Thr Thr Leu Lys Pro Ile Val Leu Ser Ile Leu Leu Ile
 1 5 10 15

Asn Thr Pro Leu Leu Ala Gln Ala His Glu Thr Glu Gln Ser Val Asp
 20 25 30

Leu Glu Thr Val Ser Val Val Gly Lys Ser Arg Pro Arg Ala Thr Ser
 35 40 45

Gly Leu Leu His Thr Ser Thr Ala Ser Asp Lys Ile Ile Ser Gly Asp
 50 55 60

Thr Leu Arg Gln Lys Ala Val Asn Leu Gly Asp Ala Leu Asp Gly Val
 65 70 75 80

Pro Gly Ile His Ala Ser Gln Tyr Gly Gly Gly Ala Ser Ala Pro Val
 85 90 95

Ile Arg Gly Gln Thr Gly Arg Arg Ile Lys Val Leu Asn His His Gly
 100 105 110

Glu Thr Gly Asp Met Ala Asp Phe Ser Pro Asp His Ala Ile Met Val
 115 120 125

Asp Thr Ala Leu Ser Gln Gln Val Glu Ile Leu Arg Gly Pro Val Thr
 130 135 140

Leu Leu Tyr Ser Ser Gly Asn Val Ala Gly Leu Val Asp Val Ala Asp
 145 150 155 160

Gly Lys Ile Pro Glu Lys Met Pro Glu Asn Gly Val Ser Gly Glu Leu
 165 170 175

Gly Leu Arg Leu Ser Ser Gly Asn Leu Glu Lys Leu Thr Ser Gly Gly
 180 185 190

Ile Asn Ile Gly Leu Gly Lys Asn Phe Val Leu His Thr Glu Gly Leu
 195 200 205

Tyr Arg Lys Ser Gly Asp Tyr Ala Val Pro Arg Tyr Arg Asn Leu Lys
 210 215 220

Arg Leu Pro Asp Ser His Ala Asp Ser Gln Thr Gly Ser Ile Gly Leu
 225 230 235 240

Ser Trp Val Gly Glu Lys Gly Phe Ile Gly Val Ala Tyr Ser Asp Arg
 245 250 255

Arg Asp Gln Tyr Gly Leu Pro Ala His Ser His Glu Tyr Asp Asp Cys
 260 265 270

His Ala Asp Ile Ile Trp Gln Lys Ser Leu Ile Asn Lys Arg Tyr Leu
 275 280 285

Gln Leu Tyr Pro His Leu Leu Thr Glu Glu Asp Ile Asp Tyr Asp Asn
 290 295 300

Pro Gly Leu Ser Cys Gly Phe His Asp Asp Asp Asn Ala His Ala His
 305 310 315 320

Thr His Ser Gly Arg Pro Trp Ile Asp Leu Arg Asn Lys Arg Tyr Glu
 325 330 335

Leu Arg Ala Glu Trp Lys Gln Pro Phe Pro Gly Phe Glu Ala Leu Arg
 340 345 350

Val His Leu Asn Arg Asn Asp Tyr Arg His Asp Glu Lys Ala Gly Asp
 355 360 365

Ala Val Glu Asn Phe Phe Asn Asn Gln Thr Gln Asn Ala Arg Ile Glu
 370 375 380

Leu Arg His Gln Pro Ile Gly Arg Leu Lys Gly Ser Trp Gly Val Gln
 385 390 395 400

Tyr Leu Gln Gln Lys Ser Ser Ala Leu Ser Ala Ile Ser Glu Ala Val
 405 410 415

Lys Gln Pro Met Leu Leu Asp Asn Lys Val Gln His Tyr Ser Phe Phe
 420 425 430

Gly Val Glu Gln Ala Asn Trp Asp Asn Phe Thr Leu Glu Gly Gly Val
 435 440 445

Arg Val Glu Lys Gln Lys Ala Ser Ile Gln Tyr Asp Lys Ala Leu Ile
 450 455 460

Asp Arg Glu Asn Tyr Tyr Asn His Pro Leu Pro Asp Leu Gly Ala His
 465 470 475 480

Arg Gln Thr Ala Arg Ser Phe Ala Leu Ser Gly Asn Trp Tyr Phe Thr
 485 490 495

Pro Gln His Lys Leu Ser Leu Thr Ala Ser His Gln Glu Arg Leu Pro
 500 505 510

Ser Thr Gln Glu Leu Tyr Ala His Gly Lys His Val Ala Thr Asn Thr
 515 520 525

Phe Glu Val Gly Asn Lys His Leu Asn Lys Glu Arg Ser Asn Asn Ile
 530 535 540

Glu Leu Ala Leu Gly Tyr Glu Gly Asp Arg Trp Gln Tyr Asn Leu Ala
 545 550 555 560

Leu Tyr Arg Asn Arg Phe Gly Asn Tyr Ile Tyr Ala Gln Thr Leu Asn
 565 570 575

Asp Gly Arg Gly Pro Lys Ser Ile Glu Asp Asp Ser Glu Met Lys Leu
 580 585 590

Val Arg Tyr Asn Gln Ser Gly Ala Asp Phe Tyr Gly Ala Glu Gly Glu
 595 600 605

Ile Tyr Phe Lys Pro Thr Pro Arg Tyr Arg Ile Gly Val Ser Gly Asp
 610 615 620

Tyr Val Arg Gly Arg Leu Lys Asn Leu Pro Ser Leu Pro Gly Arg Glu
625 630 635 640

Asp Ala Tyr Gly Asn Arg Pro Phe Ile Ala Gln Asp Asp Gln Asn Ala
645 650 655

Pro Arg Val Pro Ala Ala Arg Leu Gly Phe His Leu Lys Ala Ser Leu
660 665 670

Thr Asp Arg Ile Asp Ala Asn Leu Asp Tyr Tyr Arg Val Phe Ala Gln
675 680 685

Asn Lys Leu Ala Arg Tyr Glu Thr Arg Thr Pro Gly His His Met Leu
690 695 700

Asn Leu Gly Ala Asn Tyr Arg Arg Asn Thr Arg Tyr Gly Glu Trp Asn
705 710 715 720

Trp Tyr Val Lys Ala Asp Asn Leu Leu Asn Gln Ser Val Tyr Ala His
725 730 735

Ser Ser Phe Leu Ser Asp Thr Pro Gln Met Gly Arg Ser Phe Thr Gly
740 745 750

Gly Val Asn Val Lys Phe
755

<210> 203

<211> 225

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(225)

<400> 203

atg agg ctg gca acc aag gat ttg gcg gaa gcc att agg aaa gga cag 48
Met Arg Leu Ala Thr Lys Asp Leu Ala Glu Ala Ile Arg Lys Gly Gln
1 5 10 15

gtt cgc aaa tca agc ttt aac aca gaa caa tta agg gca att gaa aaa 96
Val Arg Lys Ser Ser Phe Asn Thr Glu Gln Leu Arg Ala Ile Glu Lys
20 25 30

gga gaa tct aaa ata ccg gat tac act tgg cat cat cat caa gat aca 144
 Gly Glu Ser Lys Ile Pro Asp Tyr Thr Trp His His His Gln Asp Thr
 35 40 45

gga agg atg caa ttg att cgt gaa ggc ttg cat cat gat acc ggc cat 192
 Gly Arg Met Gln Leu Ile Arg Glu Gly Leu His His Asp Thr Gly His
 50 55 60

att ggt tgg gaa gca atg aac aaa gga agg taa 225
 Ile Gly Trp Glu Ala Met Asn Lys Gly Arg
 65 70 75

<210> 204

<211> 74

<212> PRT

<213> Neisseria meningitidis

<400> 204

Met Arg Leu Ala Thr Lys Asp Leu Ala Glu Ala Ile Arg Lys Gly Gln
 1 5 10 15

Val Arg Lys Ser Ser Phe Asn Thr Glu Gln Leu Arg Ala Ile Glu Lys
 20 25 30

Gly Glu Ser Lys Ile Pro Asp Tyr Thr Trp His His His Gln Asp Thr
 35 40 45

Gly Arg Met Gln Leu Ile Arg Glu Gly Leu His His Asp Thr Gly His
 50 55 60

Ile Gly Trp Glu Ala Met Asn Lys Gly Arg
 65 70

<210> 205

<211> 1176

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(1176)

<400> 205

atg gca aaa atg atg aaa tgg gcg gct gtt gcg gcg gtc gcg gcg gca 48
 Met Ala Lys Met Met Lys Trp Ala Ala Val Ala Ala Val Ala Ala Ala

1	5	10	15	
gcg gtt tgg ggc gga tgg tct tat ctg aag ccc gag ccg cag gct gct				96
Ala Val Trp Gly Gly Trp Ser Tyr Leu Lys Pro Glu Pro Gln Ala Ala				
20		25	30	
tat att acg gaa acg gtc agg cgc ggc gac atc agc cgg acg gtt tct				144
Tyr Ile Thr Glu Thr Val Arg Arg Gly Asp Ile Ser Arg Thr Val Ser				
35		40	45	
gca aca ggg gag att tcg ccg tcc aac ctg gta tcg gtc ggc gcg cag				192
Ala Thr Gly Glu Ile Ser Pro Ser Asn Leu Val Ser Val Gly Ala Gln				
50		55	60	
gca tcg ggg cag att aag aaa ctt tat gtc aaa ctc ggg caa cag gtt				240
Ala Ser Gly Gln Ile Lys Lys Leu Tyr Val Lys Leu Gly Gln Gln Val				
65		70	75	80
aaa aag ggc gat ttg att gcg gaa atc aat tcg acc tcg cag acc aat				288
Lys Lys Gly Asp Leu Ile Ala Glu Ile Asn Ser Thr Ser Gln Thr Asn				
85		90	95	
acg ctc aat acg gaa aaa tcc aaa ttg gaa acg tat cag gcg aag ctg				336
Thr Leu Asn Thr Glu Lys Ser Lys Leu Glu Thr Tyr Gln Ala Lys Leu				
100		105	110	
gtg tcg gca cag att gca ttg ggc agc gcg gag aag aaa tat aag cgt				384
Val Ser Ala Gln Ile Ala Leu Gly Ser Ala Glu Lys Lys Tyr Lys Arg				
115		120	125	
cag gcg gcg ttg tgg aag gat gat gcg acc gct aaa gaa gat ttg gaa				432
Gln Ala Ala Leu Trp Lys Asp Asp Ala Thr Ala Lys Glu Asp Leu Glu				
130		135	140	
agc gca cag gat gcg ctt gcc gcc gcc aaa gcc aat gtt gcc gag ctg				480
Ser Ala Gln Asp Ala Leu Ala Ala Ala Lys Ala Asn Val Ala Glu Leu				
145		150	155	160
aag gct cta atc aga cag agc aaa att tcc atc aat acc gcc gag tcg				528
Lys Ala Leu Ile Arg Gln Ser Lys Ile Ser Ile Asn Thr Ala Glu Ser				
165		170	175	
gaa ttg ggc tac acg cgc att acc gca acg atg gac ggc acg gtg gtg				576
Glu Leu Gly Tyr Thr Arg Ile Thr Ala Thr Met Asp Gly Thr Val Val				
180		185	190	
gcg att ctc gtg gaa gag ggg cag act gtg aac gcg gcg cag tct acg				624
Ala Ile Leu Val Glu Glu Gly Gln Thr Val Asn Ala Ala Gln Ser Thr				

195	200	205	
ccg acg att gtc caa ttg gcg aat ctg gat atg atg ttg aac aaa atg			672
Pro Thr Ile Val Gln Leu Ala Asn Leu Asp Met Met Leu Asn Lys Met			
210	215	220	
cag att gcc gag ggc gat att acc aag gtg aag gcg ggg cag gat att			720
Gln Ile Ala Glu Gly Asp Ile Thr Lys Val Lys Ala Gly Gln Asp Ile			
225	230	235	240
tcg ttt acg att ttg tcc gaa ccg gat acg ccg att aag gcg aag ctc			768
Ser Phe Thr Ile Leu Ser Glu Pro Asp Thr Pro Ile Lys Ala Lys Leu			
245	250	255	
gac agc gtc gac ccc ggg ctg acc acg atg tcg tcg ggc ggc tac aac			816
Asp Ser Val Asp Pro Gly Leu Thr Thr Met Ser Ser Gly Gly Tyr Asn			
260	265	270	
agc agt acg gat acg gct tcc aat gcg gtc tac tat tat gcc cgt tcg			864
Ser Ser Thr Asp Thr Ala Ser Asn Ala Val Tyr Tyr Tyr Ala Arg Ser			
275	280	285	
ttt gtg ccg aat ccg gac ggc aaa ctc gcc acg ggg atg acg acg cag			912
Phe Val Pro Asn Pro Asp Gly Lys Leu Ala Thr Gly Met Thr Thr Gln			
290	295	300	
aat acg gtt gaa atc gac ggt gtg aaa aat gtg ctg att att ccg tcg			960
Asn Thr Val Glu Ile Asp Gly Val Lys Asn Val Leu Ile Ile Pro Ser			
305	310	315	320
ctg acc gtg aaa aat cgc ggc ggc agg gcg ttt gtg cgc gtg ttg ggt			1008
Leu Thr Val Lys Asn Arg Gly Gly Arg Ala Phe Val Arg Val Leu Gly			
325	330	335	
gca gac ggc aag gcg gcg gaa cgc gaa atc cgg acc ggt atg aga gac			1056
Ala Asp Gly Lys Ala Ala Glu Arg Glu Ile Arg Thr Gly Met Arg Asp			
340	345	350	
agt atg aat acc gaa gta aaa agc ggg ttg aaa gag ggg gac aaa gtg			1104
Ser Met Asn Thr Glu Val Lys Ser Gly Leu Lys Glu Gly Asp Lys Val			
355	360	365	
gtc atc tcc gaa ata acc gcc gcc gag cag cag gaa agc ggc gaa cgc			1152
Val Ile Ser Glu Ile Thr Ala Ala Glu Gln Gln Glu Ser Gly Glu Arg			
370	375	380	
gcc cta ggc ggc ccg ccg cgc cga			1176
Ala Leu Gly Gly Pro Pro Arg Arg			

385

390

<210> 206

<211> 392

<212> PRT

<213> Neisseria meningitidis

<400> 206

Met Ala Lys Met Met Lys Trp Ala Ala Val Ala Ala Val Ala Ala Ala

1 5 10 15

Ala Val Trp Gly Gly Trp Ser Tyr Leu Lys Pro Glu Pro Gln Ala Ala

20 25 30

Tyr Ile Thr Glu Thr Val Arg Arg Gly Asp Ile Ser Arg Thr Val Ser

35 40 45

Ala Thr Gly Glu Ile Ser Pro Ser Asn Leu Val Ser Val Gly Ala Gln

50 55 60

Ala Ser Gly Gln Ile Lys Lys Leu Tyr Val Lys Leu Gly Gln Gln Val

65 70 75 80

Lys Lys Gly Asp Leu Ile Ala Glu Ile Asn Ser Thr Ser Gln Thr Asn

85 90 95

Thr Leu Asn Thr Glu Lys Ser Lys Leu Glu Thr Tyr Gln Ala Lys Leu

100 105 110

Val Ser Ala Gln Ile Ala Leu Gly Ser Ala Glu Lys Lys Tyr Lys Arg

115 120 125

Gln Ala Ala Leu Trp Lys Asp Asp Ala Thr Ala Lys Glu Asp Leu Glu

130 135 140

Ser Ala Gln Asp Ala Leu Ala Ala Ala Lys Ala Asn Val Ala Glu Leu

145 150 155 160

Lys Ala Leu Ile Arg Gln Ser Lys Ile Ser Ile Asn Thr Ala Glu Ser

165 170 175

Glu Leu Gly Tyr Thr Arg Ile Thr Ala Thr Met Asp Gly Thr Val Val

180 185 190

Ala Ile Leu Val Glu Glu Gly Gln Thr Val Asn Ala Ala Gln Ser Thr

195 200 205

Pro Thr Ile Val Gln Leu Ala Asn Leu Asp Met Met Leu Asn Lys Met
 210 215 220

Gln Ile Ala Glu Gly Asp Ile Thr Lys Val Lys Ala Gly Gln Asp Ile
 225 230 235 240

Ser Phe Thr Ile Leu Ser Glu Pro Asp Thr Pro Ile Lys Ala Lys Leu
 245 250 255

Asp Ser Val Asp Pro Gly Leu Thr Thr Met Ser Ser Gly Gly Tyr Asn
 260 265 270

Ser Ser Thr Asp Thr Ala Ser Asn Ala Val Tyr Tyr Tyr Ala Arg Ser
 275 280 285

Phe Val Pro Asn Pro Asp Gly Lys Leu Ala Thr Gly Met Thr Thr Gln
 290 295 300

Asn Thr Val Glu Ile Asp Gly Val Lys Asn Val Leu Ile Ile Pro Ser
 305 310 315 320

Leu Thr Val Lys Asn Arg Gly Gly Arg Ala Phe Val Arg Val Leu Gly
 325 330 335

Ala Asp Gly Lys Ala Ala Glu Arg Glu Ile Arg Thr Gly Met Arg Asp
 340 345 350

Ser Met Asn Thr Glu Val Lys Ser Gly Leu Lys Glu Gly Asp Lys Val
 355 360 365

Val Ile Ser Glu Ile Thr Ala Ala Glu Gln Gln Glu Ser Gly Glu Arg
 370 375 380

Ala Leu Gly Gly Pro Pro Arg Arg
 385 390

<210> 207

<211> 342

<212> DNA

<213> Neisseria meningitidis

<220>

<221> CDS

<222> (1)..(342)

<400> 207


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ttg ttt gta tgc ttt gaa aaa tgt ttg ttt ccg gat ttt gcc att ccc 48
Leu Phe Val Cys Phe Glu Lys Cys Leu Phe Pro Asp Phe Ala Ile Pro
  1             5             10             15

atc cgg ttt tgc gct gta cga tgt gtt tta gcg cgg act tgc tca aaa 96
Ile Arg Phe Cys Ala Val Arg Cys Val Leu Ala Arg Thr Cys Ser Lys
          20             25             30

tcg gat gtg gtt ccg gta ttt ggg gct ttg att agg gat gcg gac ttt 144
Ser Asp Val Val Pro Val Phe Gly Ala Leu Ile Arg Asp Ala Asp Phe
      35             40             45

caa tat att ttt ctc agc tac aac aac gaa ggc ttg atg tct gtc ggg 192
Gln Tyr Ile Phe Leu Ser Tyr Asn Asn Glu Gly Leu Met Ser Val Gly
      50             55             60

cag gta agg gag att ttt gag cgt ttc ggc aaa tat aat ttg gtt caa 240
Gln Val Arg Glu Ile Phe Glu Arg Phe Gly Lys Tyr Asn Leu Val Gln
      65             70             75             80

acg gaa tat cgg cgt ttt aag gca gat aag aca gaa aac cgt aat cat 288
Thr Glu Tyr Arg Arg Phe Lys Ala Asp Lys Thr Glu Asn Arg Asn His
          85             90             95

aag gca aat tcg ata ttc gaa tat ctg cat gag acc ttt gca aaa ata 336
Lys Ala Asn Ser Ile Phe Glu Tyr Leu His Glu Thr Phe Ala Lys Ile
      100             105             110

gtc tgt 342
Val Cys

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<210> 208

<211> 114

<212> PRT

<213> *Neisseria meningitidis*

<400> 208

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Leu Phe Val Cys Phe Glu Lys Cys Leu Phe Pro Asp Phe Ala Ile Pro
  1             5             10             15

Ile Arg Phe Cys Ala Val Arg Cys Val Leu Ala Arg Thr Cys Ser Lys
          20             25             30

Ser Asp Val Val Pro Val Phe Gly Ala Leu Ile Arg Asp Ala Asp Phe
      35             40             45

Gln Tyr Ile Phe Leu Ser Tyr Asn Asn Glu Gly Leu Met Ser Val Gly

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50 55 60
 Gln Val Arg Glu Ile Phe Glu Arg Phe Gly Lys Tyr Asn Leu Val Gln
 65 70 75 80
 Thr Glu Tyr Arg Arg Phe Lys Ala Asp Lys Thr Glu Asn Arg Asn His
 85 90 95
 Lys Ala Asn Ser Ile Phe Glu Tyr Leu His Glu Thr Phe Ala Lys Ile
 100 105 110
 Val Cys

<210> 209
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:Primer Sequence

<400> 209
 ttggttaatt ggttgtaaca ctgg 24

<210> 210
 <211> 19
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:Primer Sequence

<400> 210
 attctcatgt ttgacagcg 19

<210> 211
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:Primer Sequence

<400> 211
atcctacaac ctcaagct 18

<210> 212
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Primer Sequence

<400> 212
atcccattct aaccaagc 18

<210> 213
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Primer Sequence

<400> 213
aagagattac gcgcagacc 19

<210> 214
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Primer Sequence

<400> 214
aatacgcaac cgcctctc 18